```
Qy    1261 ACCATCCAGCCGGACCTCAGCACCACCACCACCACCTACCAGGGCAGTCTCTGTCCCCGG 1320
           | |||||||||| ||||||||||||||||||| ||||||||||||||||||||| ||| |  ||
Db    1261 ACCATCCAGCCAGACCTCAGCACCACCACTACCACCTACCAGGGCAGTCTATGTTCGAGG 1320

Qy    1321 CAGGATGGGCCCAGCCCCAAGTTCCAGCTCACCAATGGGCACCTGCTCAGCCCCCTGGGT 1380
           |||||||| ||||||||||||||||||||||||||| | ||||| ||||||||||||| |||||
Db    1321 CAGGATGGACCCAGCCCCAAGTTCCAGCTCTCTAATGGTCACCTGCTCAGCCCACTGGGG 1380

Qy    1381 GGCGGCCGCCACACACTGCACCACAGCTCTCCCACCTCTGAGGCCGAGGAGTTCGTCTCC 1440
           | ||||||||| || |||||||||||| ||||||||||||| ||||| |||||||||
Db    1381 AGTGGCCGCCATACGTTGCACCACAGCTCACCCACCTCTGAGGCTGAGGACTTCGTCTCC 1440

Qy    1441 CGCCTCTCCACCCAGAACTACTTCCGCTCCCTGCCCCGAGGCACCAGCAACATGACCTAT 1500
           ||||||||||||| |||||||| || |||||||||||| ||||||||||||||||| ||||
Db    1441 CGCCTCTCCACCCAAAACTACTTTCGTTCCCTGCCCCGCGGCACCAGCAACATGGCCTAC 1500

Qy    1501 GGGACCTTCAACTTCCTCGGGGGCCGGCTGATGATCCCTAATACAGGTATCAGCCTCCTC 1560
           ||||||||||||||||||||||||||||||||||||||||||||| || |||||||||||
Db    1501 GGGACCTTCAACTTCCTCGGGGGCCGGCTGATGATCCCTAATACGGGGATCAGCCTCCTC 1560

Qy    1561 ATCCCCCCAGATGCCATACCCCGAGGGAAGATCTATGAGATCTACCTCACGCTGCACAAG 1620
           || ||||| ||||||| |||||||| ||||||| ||||||||||||||||| |||||||||
Db    1561 ATACCCCCGGATGCCATCCCCCGAGGAAAGATCTACGAGATCTACCTCACACTGCACAAG 1620

Qy    1621 CCGGAAGACGTGAGGTTGCCCCTAGCTGGCTGTCAGACCCTGCTGAGTCCCATCGTTAGC 1680
           || |||||||||||||||||||||||||||||||||||||||||||||||| ||||||||
Db    1621 CCAGAAGACGTGAGGTTGCCCCTAGCTGGCTGTCAGACCCTGCTGAGTCCAGTCGTTAGC 1680

Qy    1681 TGTGGACCCCCTGGCGTCCTGCTCACCCGGCCAGTCATCCTGGCTATGGACCACTGTGGG 1740
           ||||| |||||| || ||||||||||||||||||||||||||||| || ||||||||||||
Db    1681 TGTGGGCCCCCAGGAGTCCTGCTCACCCGGCCAGTCATCCTTGCAATGGACCACTGTGGA 1740

Qy    1741 GAGCCCAGCCCTGACAGCTGGAGCCTGCGCCTCAAAAAGCAGTCGTGCGAGGGCAGCTGG 1800
           ||||||||||||||||||||||||| |||||||||||||||||||| |||||||||| |||
Db    1741 GAGCCCAGCCCTGACAGCTGGAGTCTGCGCCTCAAAAAGCAGTCCTGCGAGGGCAGTTGG 1800

Qy    1801 GAGGATGTGCTGCACCTGGGCGAGGAGGCGCCCTCCCACCTCTACTACTGCCAGCTGGAG 1860
           ||||||||||||||||| || |||||| | || ||||||||||||||||||||||||||||
Db    1801 GAGGATGTGCTGCACCTTGGTGAGGAGTCACCTTCCCACCTCTACTACTGCCAGCTGGAG 1860

Qy    1861 GCCAGTGCCTGCTACGTCTTCACCGAGCAGCTGGGCCGCTTTGCCCTGGTGGGAGAGGCC 1920
           ||| | ||||||||| |||||||| ||||||| ||||||||||||||||||| |||||||||
Db    1861 GCCGGGGCCTGCTATGTCTTCACGGAGCAGCTGGGCCGCTTTGCCCTGGTAGGAGAGGCC 1920

Qy    1921 CTCAGCGTGGCTGCCGCCAAGCGCCTCAAGCTGCTTCTGTTTGCGCCGGTGGCCTGCACC 1980
           |||||||||||||||||| ||||||||||| ||| |||||||||| ||| |||||||| ||
Db    1921 CTCAGCGTGGCTGCCACCAAGCGCCTCAGGCTCCTTCTGTTTGCTCCCGTGGCCTGTACG 1980

Qy    1981 TCCCTCGAGTACAACATCCGGGTCTACTGCCTGCATGACACCCACGATGCACTCAAGGAG 2040
           ||||| |||||||||||||| || ||||||||| || ||||||||| || |||||||||
Db    1981 TCCCTTGAGTACAACATCCGAGTGTACTGCCTACGGACACCCACGACGCTCTCAAGGAG 2040

Qy    2041 GTGGTGCAGCTGGAGAAGCAGCTGGGGGGACAGCTGATCCAGGAGCCACGGGTCCTGCAC 2100
           ||||||||||||||||||||||||| || ||||||||||||||||||| || |||||||||
Db    2041 GTGGTGCAGCTGGAGAAGCAGCTAGGTGGACAGCTGATCCAGGAGCCTCGCGTCCTGCAC 2100

Qy    2101 TTCAAGGACAGTTACCACAACCTGCGCCTATCCATCCACGATGTGCCCAGCTCCCTGTGG 2160
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         |||||  ||||||||||||||||  ||  ||  |||||||||||  |||||||||||||||||||||
Db    2101 TTCAAAGACAGTTACCACAACCTACGTCTCTCCATCCACGACGTGCCCAGCTCCCTGTGG 2160

Qy    2161 AAGAGTAAGCTCCTTGTCAGCTACCAGGAGATCCCCTTTTATCACATCTGGAATGGCACG 2220
         |||||  |||||  ||||||||||||||||||||||||  |||||  |||||||||||  |||||
Db    2161 AAGAGCAAGCTACTTGTCAGCTACCAGGAGATCCCTTTTTACCACATCTGGAACGGCACC 2220

Qy    2221 CAGCGGTACTTGCACTGCACCTTCACCCTGGAGCGTGTCAGCCCCAGCACTAGTGACCTG 2280
         ||||  |||   |||||||||||||||||||||||||||  |||  |  |||||||  ||  ||||||
Db    2221 CAGCAGTATCTGCACTGCACCTTCACCCTGGAGCGCATCAACGCCAGCACCAGCGACCTG 2280

Qy    2281 GCCTGCAAGCTGTGGGTGTGGCAGGTGGAGGGCGACGGGCAGAGCTTCAGCATCAACTTC 2340
         ·|||||||||  ||||||||||||||||||||||||||  ||  |||||||||||||  |||||||||||
Db    2281 GCCTGCAAGGTGTGGGTGTGGCAGGTGGAGGGAGATGGGCAGAGCTTCAACATCAACTTC 2340

Qy    2341 AACATCACCAAGGACACAAGGTTTGCTGAGCTGCTGGCTCTGGAGAGTGAAGCGGGGGTC 2400
         ||||||||  ||·|||||||||||||||||  ||  |||||||||||||||||||||||  |||||||
Db    2341 AACATCACTAAGGACACAAGGTTTGCTGAATTGTTGGCTCTGGAGAGTGAAGGGGGGGTC 2400

Qy    2401 CCAGCCCTGGTGGGCCCCAGTGCCTTCAAGATCCCCTTCCTCATTCGGCAGAAGATAATT 2460
         |||||||||||||||||||||||||||||||||||||||||||||||||||||||  |||||  ||
Db    2401 CCAGCCCTGGTGGGCCCCAGTGCCTTCAAGATCCCCTTCCTCATTCGGCAAAAGATCATC 2460

Qy    2461 TCCAGCCTGGACCCACCCTGTAGGCGGGGTGCCGACTGGCGGACTCTGGCCCAGAAACTC 2520
         ||||  |||||||||||||||  |||||  |||||||||  |  |||||  |||||||||||
Db    2461 GCCAGTCTGGACCCACCCTGCAGCCGGGGCGCCGACTGGAGAACTCTAGCCCAGAAACTT 2520

Qy    2521 CACCTGGACAGCCATCTCAGCTTCTTTGCCTCCAAGCCCAGCCCCACAGCCATGATCCTC 2580
         ||||||||||||||||||  |||||||||||||||||||||||||||  |||||||||||||||
Db    2521 CACCTGGACAGCCATCTTAGCTTCTTTGCCTCCAAGCCCAGCCCTACAGCCATGATCCTC 2580

Qy    2581 AACCTGTGGGAGGCGCGGCACTTCCCCAACGGCAACCTCAGCCAGCTGGCTGCAGCAGTG 2640
         |||||  |||||||||  ||||||||||||||||||||||||||||  |||||||||  |||||  |||
Db    2581 AACCTATGGGAGGCACGGCACTTCCCCAACGGCAACCTCGGCCAGCTGGCAGCAGCTGTG 2640

Qy    2641 GCTGGACTGGGCCAGCCAGACGCTGGCCTCTTCACAGTGTCGGAGGCTGAGTGCTGA 2697
         ||  |||||||||||  |||||  |||||||||||||  ||||||||||||  |||  |||
Db    2641 GCCGGACTGGGCCAACCAGATGCTGGCCTCTTCACGGTGTCGGAGGCCGAGTGTTGA 2697
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RESULT 2
US-09-306-902A-1
; Sequence 1, Application US/09306902A
; Patent No. 6277585
;   GENERAL INFORMATION:
;   APPLICANT: Tessier-Lavigne, Marc
;   Leonardo, E. David
;   Hink, Lindsay
;   Masu, Masayuki
;   Kazuko, Keino-Masu
;   TITLE OF INVENTION: Netrin Receptors
;   NUMBER OF SEQUENCES: 9
;   CORRESPONDENCE ADDRESS:
;   ADDRESSEE: SCIENCE & TECHNOLOGY LAW GROUP
;   STREET: 268 BUSH STREET, SUITE 3200
;   CITY: SAN FRANCISCO
;   STATE: CALIFORNIA

```
;                    COUNTRY: USA
;                    ZIP: 94104
;              COMPUTER READABLE FORM:
;                    MEDIUM TYPE: Floppy disk
;                    COMPUTER: IBM PC compatible
;                    OPERATING SYSTEM: PC-DOS/MS-DOS
;                    SOFTWARE: PatentIn Release #1.0, Version #1.30
;              CURRENT APPLICATION DATA:
;                    APPLICATION NUMBER: US/09/306,902A
;                    FILING DATE: 07-May-1999
;                    CLASSIFICATION: <Unknown>
;              ATTORNEY/AGENT INFORMATION:
;                    NAME: OSMAN, RICHARD A
;                    REGISTRATION NUMBER: 36,627
;                    REFERENCE/DOCKET NUMBER: UC96-217
;              TELECOMMUNICATION INFORMATION:
;                    TELEPHONE: (415) 343-4341
;                    TELEFAX: (415) 343-4342
;      INFORMATION FOR SEQ ID NO: 1:
;              SEQUENCE CHARACTERISTICS:
;                    LENGTH: 3014 base pairs
;                    TYPE: nucleic acid
;                    STRANDEDNESS: double
;                    TOPOLOGY: linear
;              MOLECULE TYPE: cDNA
;              SEQUENCE DESCRIPTION: SEQ ID NO: 1:
US-09-306-902A-1

   Query Match           83.5%;  Score 2252.2;  DB 3;  Length 3014;
   Best Local Similarity  89.7%;  Pred. No. 0;
   Matches 2419;  Conservative   0;  Mismatches  278;  Indels   0;  Gaps    0;


Qy        1 ATGGCCGTCCGGCCCGGCCTGTGGCCAGCGCTCCTGGGCATAGTCCTCGCCGCTTGGCTC 60
            ||||||||||||||||||||||||||||||| ||||||||||||||||||||||||| |||||
Db        1 ATGGCCGTCCGGCCCGGCCTGTGGCCAGTGCTCCTGGGCATAGTCCTCGCCGCCTGGCTT 60

Qy       61 CGCGGCTCGGGTGCCCAGCAGAGTGCCACCGTGGCCAACCCAGTGCCTGGTGCCAACCCG 120
            || || |||||||||||||||||||||||||| ||||||||| ||||||||| |||||||||||
Db       61 CGTGGTTCGGGTGCCCAGCAGAGTGCCACGGTGGCCAATCCAGTGCCCGGTGCCAACCCC 120

Qy      121 GACCTGCTTCCCCACTTCCTGGTGGAGCCCGAGGATGTGTACATCGTCAAGAACAAGCCA 180
            |||||||| ||||||||||||||| ||||| ||||| ||||||||| ||||||||||||||||
Db      121 GACCTGCTGCCCCACTTCCTGGTAGAGCCTGAGGACGTGTACATTGTCAAGAACAAGCCG 180

Qy      181 GTGCTGCTTGTGTGCAAGGCCGTGCCCGCCACGCAGATCTTCTTCAAGTGCAACGGGGAG 240
            ||| || | |||||||||||| ||||| ||||| |||||||||||||||||||||||| |||||
Db      181 GTGTTGTTGGTGTGCAAGGCTGTGCCTGCCACCCAGATCTTCTTCAAGTGCAATGGGGAA 240

Qy      241 TGGGTGCGCCAGGTGGACCACGTGATCGAGCGCAGCACAGACGGGAGCAGTGGGCTGCCC 300
            ||||| ||||||||| || ||||| || || ||||||||| ||| | |||||| || ||||
Db      241 TGGGTCCGCCAGGTCGATCACGTAATTGAACGCAGCACCGACAGCAGCAGCGGATTGCCA 300

Qy      301 ACCATGGAGGTCCGCATTAATGTCTCAAGGCAGCAGGTCGAGAAGGTGTTCGGGCTGGAG 360
            |||||||||||||| || || || || |||||||||| ||||| |||||| |||||||||
Db      301 ACCATGGAGGTCCGTATCAACGTATCGAGGCAGCAGGTAGAGAAAGTGTTTGGGCTGGAG 360

Qy      361 GAATACTGGTGCCAGTGCGTGGCATGGAGCTCCTCGGGCACCACCAAGAGTCAGAAGGCC 420
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          |||||||||||||||||||| ||||||||||||||||||||||| ||||||||| |||||||||||
Db    361 GAATACTGGTGCCAGTGTGTGGCATGGAGCTCCTCGGGTACCACCAAAAGTCAGAAGGCC 420

Qy    421 TACATCCGCATAGCCAGATTGCGCAAGAACTTCGAGCAGGAGCCGCTGGCCAAGGAGGTG 480
          |||||||| || |||   |||||||||||||||| |||||||||| ||||||||||| |||
Db    421 TACATCCGGATTGCCTATTTGCGCAAGAACTTTGAGCAGGAGCCACTGGCCAAGGAAGTG 480

Qy    481 TCCCTGGAGCAGGGCATCGTGCTGCCCTGCCGTCCACCGGAGGGCATCCCTCCAGCCGAG 540
          || ||||||||| ||||| || || || || || || || || || ||||| ||||| |||
Db    481 TCACTGGAGCAAGGCATTGTACTACCTTGTCGCCCCCAGAAGGAATCCCCCCAGCTGAG 540

Qy    541 GTGGAGTGGCTCCGGAACGAGGACCTGGTGGACCCGTCCCTGGACCCCAATGTATACATC 600
          |||||||||| || || ||||||||| |||||||||| ||||| || ||||||| ||||||
Db    541 GTGGAGTGGCTTCGAAATGAGGACCTCGTGGACCCCTCCCTCGATCCCAATGTGTACATC 600

Qy    601 ACGCGGGAGCACAGCCTGGTGGTGCGACAGGCCCGCCTTGCTGACACGGCCAACTACACC 660
          |||||||||||||||||| || |||||| |||||||||| || |||||||||||||||||||
Db    601 ACGCGGGAGCACAGCCTAGTCGTGCGTCAGGCCCGCCTGGCCGACACGGCCAACTACACC 660

Qy    661 TGCGTGGCCAAGAACATCGTGGCACGTCGCCGCAGCGCCTCCGCTGCTGTCATCGTCTAC 720
          || |||||||||||||||||| || ||||||||| |||| || || ||||| || ||
Db    661 TGTGTGGCCAAGAACATCGTAGCCCGTCGCCGAAGCACCTCTGCAGCGGTCATTGTTTAT 720

Qy    721 GTGAACGGTGGGTGGTCGACGTGGACCGAGTGGTCCGTCTGCAGCGCCAGCTGTGGGCGC 780
          |||||||||||||||||||||||||||| |||||||||||||||||||||||||||||||
Db    721 GTGAACGGTGGGTGGTCGACGTGGACTGAGTGGTCCGTCTGCAGCGCCAGCTGTGGGCGT 780

Qy    781 GGCTGGCAGAAACGGAGCCGGAGCTGCACCAACCCGGCGCCTCTCAACGGGGGCGCTTTC 840
          ||||||||||||||||||||||||||||||||||||||| |||||||||||||||||| |||
Db    781 GGCTGGCAGAAACGGAGCCGGAGCTGCACCAACCCGGCACCTCTCAACGGGGGCGCCTTC 840

Qy    841 TGTGAGGGGCAGAATGTCCAGAAAACAGCCTGCGCCACCCTGTGCCCAGTAGACGGCAGC 900
          |||||||||||||||||||||||||||||||||||||||||| |||||||||| || || |||
Db    841 TGTGAGGGGCAGAATGTCCAGAAAACAGCCTGCGCCACTCTGTGCCCAGTGGATGGGAGC 900

Qy    901 TGGAGCCCGTGGAGCAAGTGGTCGGCCTGTGGGCTGGACTGCACCCACTGGCGGAGCCGT 960
          |||||   |||||||| ||||||| ||||||| |||||||||| |||||||||||||||||||
Db    901 TGGAGTTCGTGGAGTAAGTGGTCAGCCTGTGGGCTTGACTGCACCCACTGGCGGAGCCGC 960

Qy    961 GAGTGCTCTGACCCAGCACCCCGCAACGGAGGGGAGGAGTGCCAGGGCACTGACCTGGAC 1020
          ||||||||||||||||||||||||||| ||||| |||||||| | ||| ||||||||||||
Db    961 GAGTGCTCTGACCCAGCACCCCGCAATGGAGGTGAGGAGTGTCGGGGTGCTGACCTGGAC 1020

Qy   1021 ACCCGCAACTGTACCAGTGACCTCTGTGTACACAGTGCTTCTGGCCCTGAGGACGTGGCC 1080
          |||||||||||||||||||||||||||| | |||| ||||||| |||| ||||||||||||
Db   1021 ACCCGCAACTGTACCAGTGACCTCTGCCTGCACACCGCTTCTTGCCCCGAGGACGTGGCT 1080

Qy   1081 CTCTATGTGGGCCTCATCGCCGTGGCCGTCTGCCTGGTCCTGCTGCTGCTTGTCCTCATC 1140
          |||||   | ||||||   |||| ||||| || ||||||   || |||||   ||||   |||
Db   1081 CTCTACATCGGCCTTGTCGCTGTGGCTGTGTGCCTCTTCTTGCTGTTGCTGGCCCTTGGA 1140

Qy   1141 CTCGTTTATTGCCGGAAGAAGGAGGGGCTGGACTCAGATGTGGCTGACTCGTCCATTCTC 1200
          ||| ||||| || || ||||||||| |||||||||| |||||||| |||||||||||| |||
Db   1141 CTCATTTACTGTCGCAAGAAGGAAGGGCTGGACTCCGATGTGGCCGACTCGTCCATCCTC 1200

Qy   1201 ACCTCAGGCTTCCAGCCCGTCAGCATCAAGCCCAGCAAAGCAGACAACCCCCATCTGCTC 1260
          |||||   |||||||||| |||||||||||||||||||||||||||||||||||||| |||||||
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Db    1201 ACCTCGGGCTTCCAGCCTGTCAGCATCAAGCCCAGCAAAGCAGACAACCCCCACCTGCTC 1260

Qy    1261 ACCATCCAGCCGGACCTCAGCACCACCACCACCACCTACCAGGGCAGTCTCTGTCCCCGG 1320
           |||||||||| |||||||||||||||| ||||||||||||||||||||||| ||| | ||
Db    1261 ACCATCCAGCCAGACCTCAGCACCACCACTACCACCTACCAGGGCAGTCTATGTTCGAGG 1320

Qy    1321 CAGGATGGGCCCAGCCCCAAGTTCCAGCTCACCAATGGGCACCTGCTCAGCCCCCTGGGT 1380
           ||||||||| |||||||||||||||||||||||| | ||||| |||||||||||| |||||
Db    1321 CAGGATGGACCCAGCCCCAAGTTCCAGCTCTCTAATGGTCACCTGCTCAGCCCACTGGGG 1380

Qy    1381 GGCGGCCGCCACACACTGCACCACAGCTCTCCCACCTCTGAGGCCGAGGAGTTCGTCTCC 1440
            | |||||||| || |||||||||||| ||||||||||||||| ||||| |||||||||
Db    1381 AGTGGCCGCCATACGTTGCACCACAGCTCACCCACCTCTGAGGCTGAGGACTTCGTCTCC 1440

Qy    1441 CGCCTCTCCACCCAGAACTACTTCCGCTCCCTGCCCCGAGGCACCAGCAACATGACCTAT 1500
           ||||||||||||| |||||||| || |||||||||||||| ||||||||||||||| ||||
Db    1441 CGCCTCTCCACCCAAAACTACTTTCGTTCCCTGCCCCGCGGCACCAGCAACATGGCCTAC 1500

Qy    1501 GGGACCTTCAACTTCCTCGGGGGCCGGCTGATGATCCCTAATACAGGTATCAGCCTCCTC 1560
           ||||||||||||||||||||||||||||||||||||||||||||||| || |||||||||||
Db    1501 GGGACCTTCAACTTCCTCGGGGGCCGGCTGATGATCCCTAATACGGGGATCAGCCTCCTC 1560

Qy    1561 ATCCCCCCAGATGCCATACCCCGAGGGAAGATCTATGAGATCTACCTCACGCTGCACAAG 1620
           || ||||| |||||||| ||||||| |||||||| |||||| |||||||||||| |||||||
Db    1561 ATACCCCCGGATGCCATCCCCCGAGGAAAGATCTACGAGATCTACCTCACACTGCACAAG 1620

Qy    1621 CCGGAAGACGTGAGGTTGCCCCTAGCTGGCTGTCAGACCCTGCTGAGTCCCATCGTTAGC 1680
           || |||||||||||||||||||||||||||||||||||||||||||||||| ||||||||
Db    1621 CCAGAAGACGTGAGGTTGCCCCTAGCTGGCTGTCAGACCCTGCTGAGTCCAGTCGTTAGC 1680

Qy    1681 TGTGGACCCCCTGGCGTCCTGCTCACCCGGCCAGTCATCCTGGCTATGGACCACTGTGGG 1740
           ||||| |||||| || |||||||||||||||||||||||||| || ||||||||||||||
Db    1681 TGTGGGCCCCCAGGAGTCCTGCTCACCCGGCCAGTCATCCTTGCAATGGACCACTGTGGA 1740

Qy    1741 GAGCCCAGCCCTGACAGCTGGAGCCTGCGCCTCAAAAAGCAGTCGTGCGAGGGCAGCTGG 1800
           |||||||||||||||||||||||| |||||||||||||||||||| ||||||||||| |||
Db    1741 GAGCCCAGCCCTGACAGCTGGAGTCTGCGCCTCAAAAAGCAGTCCTGCGAGGGCAGTTGG 1800

Qy    1801 GAGGATGTGCTGCACCTGGGCGAGGAGGCGCCCTCCCACCTCTACTACTGCCAGCTGGAG 1860
           |||||||||||||||| || |||||| | || ||||||||||||||||||||||||||||
Db    1801 GAGGATGTGCTGCACCTTGGTGAGGAGTCACCTTCCCACCTCTACTACTGCCAGCTGGAG 1860

Qy    1861 GCCAGTGCCTGCTACGTCTTCACCGAGCAGCTGGGCCGCTTTGCCCTGGTGGGAGAGGCC 1920
           ||| | ||||||||| |||||||||||||||||||||||||||||||||||| |||||||||
Db    1861 GCCGGGGCCTGCTATGTCTTCACGGAGCAGCTGGGCCGCTTTGCCCTGGTAGGAGAGGCC 1920

Qy    1921 CTCAGCGTGGCTGCCGCCAAGCGCCTCAAGCTGCTTCTGTTTGCGCCGGTGGCCTGCACC 1980
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Db    1921 CTCAGCGTGGCTGCCACCAAGCGCCTCAGGCTCCTTCTGTTTGCTCCCGTGGCCTGTACG 1980

Qy    1981 TCCCTCGAGTACAACATCCGGGTCTACTGCCTGCATGACACCCACGATGCACTCAAGGAG 2040
           ||||| ||||||||||||||| || |||||||| || |||||||||| || |||||||||
Db    1981 TCCCTTGAGTACAACATCCGAGTGTACTGCCTACACGACACCCACGACGCTCTCAAGGAG 2040

Qy    2041 GTGGTGCAGCTGGAGAAGCAGCTGGGGGGACAGCTGATCCAGGAGCCACGGGTCCTGCAC 2100
           |||||||||||||||||||||||||| || |||||||||||||||||||| || ||||||||
Db    2041 GTGGTGCAGCTGGAGAAGCAGCTAGGTGGACAGCTGATCCAGGAGCCTCGCGTCCTGCAC 2100
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Qy      2101 TTCAAGGACAGTTACCACAACCTGCGCCTATCCATCCACGATGTGCCCAGCTCCCTGTGG 2160
             ||||| ||||||||||||||||| || || |||||||||||| ||||||||||||||||||
Db      2101 TTCAAAGACAGTTACCACAACCTACGTCTCTCCATCCACGACGTGCCCAGCTCCCTGTGG 2160

Qy      2161 AAGAGTAAGCTCCTTGTCAGCTACCAGGAGATCCCCTTTTATCACATCTGGAATGGCACG 2220
             ||||| ||||| ||||||||||||||||||||||||||| ||||| |||||||||| |||||
Db      2161 AAGAGCAAGCTACTTGTCAGCTACCAGGAGATCCCTTTTTACCACATCTGGAACGGCACC 2220

Qy      2221 CAGCGGTACTTGCACTGCACCTTCACCCTGGAGCGTGTCAGCCCCAGCACTAGTGACCTG 2280
             |||| |||  |||||||||||||||||||||||||| ||| | ||||||| || ||||||
Db      2221 CAGCAGTATCTGCACTGCACCTTCACCCTGGAGCGCATCAACGCCAGCACCAGCGACCTG 2280

Qy      2281 GCCTGCAAGCTGTGGGTGTGGCAGGTGGAGGGCGACGGGCAGAGCTTCAGCATCAACTTC 2340
             |||||||| |||||||||||||||||||||||| || |||||||||||| |||||||||||
Db      2281 GCCTGCAAGGTGTGGGTGTGGCAGGTGGAGGGAGATGGGCAGAGCTTCAACATCAACTTC 2340

Qy      2341 AACATCACCAAGGACACAAGGTTTGCTGAGCTGCTGGCTCTGGAGAGTGAAGCGGGGGTC 2400
             |||||||| ||||||||||||||||||||  || ||||||||||||||||||| |||||||
Db      2341 AACATCACTAAGGACACAAGGTTTGCTGAATTGTTGGCTCTGGAGAGTGAAGGGGGGGTC 2400

Qy      2401 CCAGCCCTGGTGGGCCCCAGTGCCTTCAAGATCCCCTTCCTCATTCGGCAGAAGATAATT 2460
             |||||||||||||||||||||||||||||||||||||||||||||||||||| ||||| ||
Db      2401 CCAGCCCTGGTGGGCCCCAGTGCCTTCAAGATCCCCTTCCTCATTCGGCAAAAGATCATC 2460

Qy      2461 TCCAGCCTGGACCCACCCTGTAGGCGGGGTGCCGACTGGCGGACTCTGGCCCAGAAACTC 2520
             |||| |||||||||||||||| || ||||| ||||||||| | ||||| |||||||||||
Db      2461 GCCAGTCTGGACCCACCCTGCAGCCGGGGCGCCGACTGGAGAACTCTAGCCCAGAAACTT 2520

Qy      2521 CACCTGGACAGCCATCTCAGCTTCTTTGCCTCCAAGCCCAGCCCCACAGCCATGATCCTC 2580
             ||||||||||||||||||||| |||||||||||||||||||||| ||||||||||||||||
Db      2521 CACCTGGACAGCCATCTTAGCTTCTTTGCCTCCAAGCCCAGCCCTACAGCCATGATCCTC 2580

Qy      2581 AACCTGTGGGAGGCGCGGCACTTCCCCAACGGCAACCTCAGCCAGCTGGCTGCAGCAGTG 2640
             ||||| |||||||| |||||||||||||||||||||||| |||||||||||||| ||| |||
Db      2581 AACCTATGGGAGGCACGGCACTTCCCCAACGGCAACCTCGGCCAGCTGGCAGCAGCTGTG 2640

Qy      2641 GCTGGACTGGGCCAGCCAGACGCTGGCCTCTTCACAGTGTCGGAGGCTGAGTGCTGA 2697
             || |||||||||| ||||| ||||||||||||||| ||||||||||| ||||| |||
Db      2641 GCCGGACTGGGCCAACCAGATGCTGGCCTCTTCACGGTGTCGGAGGCCGAGTGTTGA 2697
```

RESULT 3
US-08-808-982-2
; Sequence 2, Application US/08808982
; Patent No. 5939271
;   GENERAL INFORMATION:
;   APPLICANT: Tessier-Lavigne, Marc
;   APPLICANT: Leonardo, E. David
;   APPLICANT: Hink, Lindsay
;   APPLICANT: Masu, Masayuki
;   APPLICANT: Kazuko, Keino-Masu
;   TITLE OF INVENTION: Netrin Receptors
;   NUMBER OF SEQUENCES: 8
;   CORRESPONDENCE ADDRESS:
;   ADDRESSEE: SCIENCE & TECHNOLOGY LAW GROUP
;   STREET: 268 BUSH STREET, SUITE 3200

; INFORMATION FOR SEQ ID NO: 2:
;   SEQUENCE CHARACTERISTICS:
;   LENGTH: 1787 base pairs
;   TYPE: nucleic acid
;   STRANDEDNESS: double
;   TOPOLOGY: linear
;   MOLECULE TYPE: cDNA
US-08-808-982-2

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  Query Match              57.6%;  Score 1552.4;  DB 2;  Length 1787;
  Best Local Similarity    98.5%;  Pred. No. 0;
  Matches 1651;  Conservative    0;  Mismatches   16;  Indels    9;  Gaps      8;

Qy      1025 GCAACTGTACCAGTGACCTCTGTGTACACAGTGCTTCTGGCCCTGAGGACGTGGCCCTCT 1084
             |||||||||||||||||||||||| ||||||| |||||||||||||||||||||||||||
Db         1 GCAACTGTACCAGTGACCTCTG-GTACACACTGCTTCTGGCCCTGAGGACGTGGCCCTCT 59

Qy      1085 ATGTGGGCCTCATCGCCGTGGCCGTCTGCCTGGTCCTGCTGCTGCTTGTCCTCATCCTCG 1144
             ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        60 ATGTGGGCCTCATCGCCGTGGCCGTCTGCCTGGTCCTGCTGCTGCTTGTCCTCATCCTCG 119

Qy      1145 TTTATTGCCGGAAGAAGGAGGGGCTGGACTCAGATGTGGCTGACTCGTCCATTCTCACCT 1204
             ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db       120 TTTATTGCCGGAAGAAGGAGGGGCTGGACTCAGATGTGGCTGACTCGTCCATTCTCACCT 179

Qy      1205 CAGGCTTCCAGCCCGTCAGCATC-AAGCCCAGCAAAGCAGACAACCCCCATCTGCTCACC 1263
             |||||||||||||||||||||||| |||||||||||||||||||||||||||||||||||
Db       180 CAGGCTTCCAGCCCGTCAGCATCTAAGCCCAGCAAAGCAGACAACCCCCATCTGCTCACC 239

Qy      1264 ATCCAGCCGGACCTCAGCACCACCACCACCACCTACCAGGGCAGTCTCTGTCCCCGGCAG 1323
             ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db       240 ATCCAGCCGGACCTCAGCACCACCACCACCACCTACCAGGGCAGTCTCTGTCCCCGGCAG 299

Qy      1324 GATGGGCCCAGCCCCAAGTTCCAGCTCACCAATGGGCACCTGCTCAGCCCCCTGGGTGGC 1383
             ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db       300 GATGGGCCCAGCCCCAAGTTCCAGCTCACCAATGGGCACCTGCTCAGCCCCCTGGGTGGC 359
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Qy  1384 GGCCGCCACACACTGCACCACAGCTCTCCCACCTCTGAGGCCGAGGAGTTCGTCTCCCGC 1443
         |||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db   360 GGCCGCCACACACTGCACCACAGCTCTCCCACCTCTGAGGCCGAGGAGTTCGTCTCCCGC 419

Qy  1444 CTCTCCACCCAGAACTACTTCCGCTCCCTGCCCCGAGGCACCAGCAACATGACCTATGGG 1503
         |||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db   420 CTCTCCACCCAGAACTACTTCCGCTCCCTGCCCCGAGGCACCAGCAACATGACCTATGGG 479

Qy  1504 ACCTTCAACTTCCTCGGGGGCCGGCTGATGATCCCTAATACAGGTATCAGCCTCCTCATC 1563
         |||||||||||||||||||||||||||||||||||||||||||||| ||||||||||||||
Db   480 ACCTTCAACTTCCTCGGGGGCCGGCTGATGATCCCTAATACAGGAATCAGCCTCCTCATC 539

Qy  1564 CCCCCAGATGCCATACCCCGAGGGAAGATCTATGAGATCTACCTCACGCTGCACAAGCCG 1623
         |||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db   540 CCCCCAGATGCCATACCCCGAGGGAAGATCTATGAGATCTACCTCACGCTGCACAAGCCG 599

Qy  1624 GAAGACGTGAGGTTGCCCCTAGCTGGCTGTCAGACCCTGCTGAGTCCCATCGTTAGCTGT 1683
         |||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db   600 GAAGACGTGAGGTTGCCCCTAGCTGGCTGTCAGACCCTGCTGAGTCCCATCGTTAGCTGT 659

Qy  1684 GGACCCCCTGGCGTCCTGCTCACCCGGCCAGTCATCCTGGCTATGGACCACTGTGGGGAG 1743
         |||||.|||||||||||||||||||||||||||||||||||||||||||||||||||||
Db   660 GGACCCCCTGGCGTCCTGCTCACCCGGCCAGTCATCCTGGCTATGGACCACTGTGGGGAG 719

Qy  1744 CCCAGCCCTGACAGCTGGAGCCTGCGCCTCAAAAAGCAGTCGTGCGAGGGCAGCTGGGAG 1803
         ||||||||||||||||||||||||    ||||||||||||||||||||||| ||||||||
Db   720 CCCAGCCCTGACAGCTGGAGCCTGGCCCTCAAAAAGCAGTCGTGCGAGGG-AGCTGGGAG 778

Qy  1804 GATGTGCTGCACCTGGGCGAGGAGGCGCCCTCCCACCTCTACTACTGCCAGCTGGAGGCC 1863
         |||||  |||||||||||||||||||||||||||||||||||||||||||||||||||||
Db   779 GATGT-CTGCACCTGGGCGAGGAGGCGCCCTCCCACCTCTACTACTGCCAGCTGGAGGCC 837

Qy  1864 AGTGCCTGCTACGTCTTCACCGAGCAGCTGGGCCGCTTTGCCCTGGTGGGAGAGGCCCTC 1923
         |||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db   838 AGTGCCTGCTACGTCTTCACCGAGCAGCTGGGCCGCTTTGCCCTGGTGGGAGAGGCCCTC 897

Qy  1924 AGCGTGGCTGCCGCCAAGCGCCTCAAGCTGCTTCTGTTTGCGCCGGTGGCCTGCACCTCC 1983
         |||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db   898 AGCGTGGCTGCCGCCAAGCGCCTCAAGCTGCTTCTGTTTGCGCCGGTGGCCTGCACCTCC 957

Qy  1984 CTCGAGTACAACATCCGGGTCTACTGCCTGCATGACACCCACGATGCACTCAAGGAGGTG 2043
         |||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db   958 CTCGAGTACAACATCCGGGTCTACTGCCTGCATGACACCCACGATGCACTCAAGGAGGTG 1017

Qy  2044 GTGCAGCTGGAGAAGCAGCTGGGGGGACAGCTGATCCAGGAGCCACGGGTCCTGCACTTC 2103
         ||||.||||||||||||||||||||||||||||||||||||||||||||||||||||.|||
Db  1018 GTGCAGCTGGAGAAGCAGCTGGGGGGACAGCTGATCCAGGAGCCACGGGTCCTGCACTT- 1076

Qy  2104 AAGGACAGTTACCACAACCTGCGCCTATCCATCCACGATGTGCCCAGCTCCCTGTGGAAG 2163
         |||||||||||||||||||||   |||    ||||||||||||||||||||||||||||||
Db  1077 AAGGACAGTTACCACAACCT--GCCCTATCATCCACGATGTGCCCAGCTCCCTGTGGAAG 1134

Qy  2164 AGTAAGCTCCTTGTCAGCTACCAGGAGATCCCCTTTTATCACATCTGGAATGGCACGCAG 2223
         |||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db  1135 AGTAAGCTCCTTGTCAGCTACCAGGAGATCCCCTTTTATCACATCTGGAATGGCACGCAG 1194

Qy  2224 CGGTACTTGCACTGCACCTTCACCCTGGAGCGTGTCAGCCCCAGCACTAGTGACCTGGCC 2283
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                 |||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db      1195  CGGTACTTGCACTGCACCTTCACCCTGGAGCGTGTCAGCCCCAGCACTAGTGACCTGGCC 1254

Qy      2284  TGCAAGCTGTGGGTGTGGCAGGTGGAGGGCGACGGGCAGAGCTTCAGCATCAACTTCAAC 2343
                 |||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db      1255  TGCAAGCTGTGGGTGTGGCAGGTGGAGGGCGACGGGCAGAGCTTCAGCATCAACTTCAAC 1314

Qy      2344  ATCACCAAGGACACAAGGTTTGCTGAGCTGCTGGCTCTGGAGAGTGAAGCGGGGGTCCCA 2403
                 |||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db      1315  ATCACCAAGGACACAAGGTTTGCTGAGCTGCTGGCTCTGGAGAGTGAAGCGGGGGTCCCA 1374

Qy      2404  GCCCTGGTGGGCCCCAGTGCCTTCAAGATCCCCTTCCTCATTCGGCAGAAGATAATTTCC 2463
                 |||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db      1375  GCCCTGGTGGGCCCCAGTGCCTTCAAGATCCCCTTCCTCATTCGGCAGAAGATAATTTCC 1434

Qy      2464  AGCCTGGACCCACCCTGTAGGCGGGGTGCCGACTGGCGGACTCTGGCCCAGAAACTCCAC 2523
                 |||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db      1435  AGCCTGGACCCACCCTGTAGGCGGGGTGCCGACTGGCGGACTCTGGCCCAGAAACTCCAC 1494

Qy      2524  CTGGACAGCCATCTCAGCTTCTTTGCCTCCAAGCCCAGCCCCACAGCCATGATCCTCAAC 2583
                 |||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db      1495  CTGGACAGCCATCTCAGCTTCTTTGCCTCCAAGCCCAGCCCCACAGCCATGATCCTCAAC 1554

Qy      2584  CTGTGGGAGGCGCGGCACTTCCCCAACGGCAACCTCAGCCAGCTGGCTGCAGCAGTGGCT 2643
                 |||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db      1555  CTGTGGGAGGCGCGGCACTTCCCCAACGGCAACCTCAGCCAGCTGGCTGCAGCAGTGGCT 1614

Qy      2644  GGACTGGGCCAGCCAGACGCTGGCCTC-TTCACAGTG-TCGGAGGCTGAGTGCTGA 2697
                 ||      |||||||  ||||  |||| || ||||||||| |||||||||||||||||
Db      1615  GGGACTGGCCAGCAGGACGGTGGCTTCTTTCACAGTGTTCGGAGGCTGAGTGCTGA 1670
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RESULT 4
US-09-306-902A-2
; Sequence 2, Application US/09306902A
; Patent No. 6277585
;   GENERAL INFORMATION:
;   APPLICANT: Tessier-Lavigne, Marc
;   Leonardo, E. David
;   Hink, Lindsay
;   Masu, Masayuki
;   Kazuko, Keino-Masu
;   TITLE OF INVENTION: Netrin Receptors
;   NUMBER OF SEQUENCES: 9
;   CORRESPONDENCE ADDRESS:
;   ADDRESSEE: SCIENCE & TECHNOLOGY LAW GROUP
;   STREET: 268 BUSH STREET, SUITE 3200
;   CITY: SAN FRANCISCO
;   STATE: CALIFORNIA
;   COUNTRY: USA
;   ZIP: 94104
;   COMPUTER READABLE FORM:
;   MEDIUM TYPE: Floppy disk
;   COMPUTER: IBM PC compatible
;   OPERATING SYSTEM: PC-DOS/MS-DOS
;   SOFTWARE: PatentIn Release #1.0, Version #1.30
;   CURRENT APPLICATION DATA:

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   Query Match               57.6%;   Score 1552.4;   DB 3;   Length 1787;
   Best Local Similarity    98.5%;   Pred. No. 0;
   Matches 1651;   Conservative    0;   Mismatches   16;   Indels   9;   Gaps     8;


Qy      1025 GCAACTGTACCAGTGACCTCTGTGTACACAGTGCTTCTGGCCCTGAGGACGTGGCCCTCT 1084
             |||||||||||||||||||||||| |||||||  |||||||||||||||||||||||||||
Db         1 GCAACTGTACCAGTGACCTCTG-GTACACACTGCTTCTGGCCCTGAGGACGTGGCCCTCT 59


Qy      1085 ATGTGGGCCTCATCGCCGTGGCCGTCTGCCTGGTCCTGCTGCTGCTTGTCCTCATCCTCG 1144
             ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        60 ATGTGGGCCTCATCGCCGTGGCCGTCTGCCTGGTCCTGCTGCTGCTTGTCCTCATCCTCG 119


Qy      1145 TTTATTGCCGGAAGAAGGAGGGGCTGGACTCAGATGTGGCTGACTCGTCCATTCTCACCT 1204
             ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db       120 TTTATTGCCGGAAGAAGGAGGGGCTGGACTCAGATGTGGCTGACTCGTCCATTCTCACCT 179


Qy      1205 CAGGCTTCCAGCCCGTCAGCATC-AAGCCCAGCAAAGCAGACAACCCCCATCTGCTCACC 1263
             |||||||||||||||||||||||| |||||||||||||||||||||||||||||||||||
Db       180 CAGGCTTCCAGCCCGTCAGCATCTAAGCCCAGCAAAGCAGACAACCCCCATCTGCTCACC 239


Qy      1264 ATCCAGCCGGACCTCAGCACCACCACCACCACCTACCAGGGCAGTCTCTGTCCCCGGCAG 1323
             |||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db       240 ATCCAGCCGGACCTCAGCACCACCACCACCACCTACCAGGGCAGTCTCTGTCCCCGGCAG 299


Qy      1324 GATGGGCCCAGCCCCAAGTTCCAGCTCACCAATGGGCACCTGCTCAGCCCCCTGGGTGGC 1383
             ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db       300 GATGGGCCCAGCCCCAAGTTCCAGCTCACCAATGGGCACCTGCTCAGCCCCCTGGGTGGC 359


Qy      1384 GGCCGCCACACACTGCACCACAGCTCTCCCACCTCTGAGGCCGAGGAGTTCGTCTCCCGC 1443
             ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db       360 GGCCGCCACACACTGCACCACAGCTCTCCCACCTCTGAGGCCGAGGAGTTCGTCTCCCGC 419


Qy      1444 CTCTCCACCCAGAACTACTTCCGCTCCCTGCCCCGAGGCACCAGCAACATGACCTATGGG 1503
             ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db       420 CTCTCCACCCAGAACTACTTCCGCTCCCTGCCCCGAGGCACCAGCAACATGACCTATGGG 479


Qy      1504 ACCTTCAACTTCCTCGGGGGCCGGCTGATGATCCCTAATACAGGTATCAGCCTCCTCATC 1563
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                   |||||||||||||||||||||||||||||||||||||||||||||||| ||||||||||||||||
Db      480 ACCTTCAACTTCCTCGGGGGCCGGCTGATGATCCCTAATACAGGAATCAGCCTCCTCATC 539        .

Qy     1564 CCCCCAGATGCCATACCCCGAGGGAAGATCTATGAGATCTACCTCACGCTGCACAAGCCG 1623
                   ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db      540 CCCCCAGATGCCATACCCCGAGGGAAGATCTATGAGATCTACCTCACGCTGCACAAGCCG 599

Qy     1624 GAAGACGTGAGGTTGCCCCTAGCTGGCTGTCAGACCCTGCTGAGTCCCATCGTTAGCTGT 1683
                   ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db      600 GAAGACGTGAGGTTGCCCCTAGCTGGCTGTCAGACCCTGCTGAGTCCCATCGTTAGCTGT 659

Qy     1684 GGACCCCCTGGCGTCCTGCTCACCCGGCCAGTCATCCTGGCTATGGACCACTGTGGGGAG 1743
                   ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db      660 GGACCCCCTGGCGTCCTGCTCACCCGGCCAGTCATCCTGGCTATGGACCACTGTGGGGAG 719

Qy     1744 CCCAGCCCTGACAGCTGGAGCCTGCGCCTCAAAAAGCAGTCGTGCGAGGGCAGCTGGGAG 1803
                   |||||||||||||||||||||||| |||||||||||||||||||||||| |||||||||
Db      720 CCCAGCCCTGACAGCTGGAGCCTGGCCCTCAAAAAGCAGTCGTGCGAGGG-AGCTGGGAG 778

Qy     1804 GATGTGCTGCACCTGGGCGAGGAGGCGCCCTCCCACCTCTACTACTGCCAGCTGGAGGCC 1863
                   ||||| ||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db      779 GATGT-CTGCACCTGGGCGAGGAGGCGCCCTCCCACCTCTACTACTGCCAGCTGGAGGCC 837

Qy     1864 AGTGCCTGCTACGTCTTCACCGAGCAGCTGGGCCGCTTTGCCCTGGTGGGAGAGGCCCTC 1923
                   ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db      838 AGTGCCTGCTACGTCTTCACCGAGCAGCTGGGCCGCTTTGCCCTGGTGGGAGAGGCCCTC 897

Qy     1924 AGCGTGGCTGCCGCCAAGCGCCTCAAGCTGCTTCTGTTTGCGCCGGTGGCCTGCACCTCC 1983
                   ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db      898 AGCGTGGCTGCCGCCAAGCGCCTCAAGCTGCTTCTGTTTGCGCCGGTGGCCTGCACCTCC 957

Qy     1984 CTCGAGTACAACATCCGGGTCTACTGCCTGCATGACACCCACGATGCACTCAAGGAGGTG 2043
                   ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db      958 CTCGAGTACAACATCCGGGTCTACTGCCTGCATGACACCCACGATGCACTCAAGGAGGTG 1017

Qy     2044 GTGCAGCTGGAGAAGCAGCTGGGGGGACAGCTGATCCAGGAGCCACGGGTCCTGCACTTC 2103
                   |||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db     1018 GTGCAGCTGGAGAAGCAGCTGGGGGGACAGCTGATCCAGGAGCCACGGGTCCTGCACTT- 1076

Qy     2104 AAGGACAGTTACCACAACCTGCGCCTATCCATCCACGATGTGCCCAGCTCCCTGTGGAAG 2163
                   ||||||||||||||||||||   |||        ||||||||||||||||||||||||||
Db     1077 AAGGACAGTTACCACAACCT--GCCCTATCATCCACGATGTGCCCAGCTCCCTGTGGAAG 1134

Qy     2164 AGTAAGCTCCTTGTCAGCTACCAGGAGATCCCCTTTTATCACATCTGGAATGGCACGCAG 2223
                   ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db     1135 AGTAAGCTCCTTGTCAGCTACCAGGAGATCCCCTTTTATCACATCTGGAATGGCACGCAG 1194

Qy     2224 CGGTACTTGCACTGCACCTTCACCCTGGAGCGTGTCAGCCCCAGCACTAGTGACCTGGCC 2283
                   |||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db     1195 CGGTACTTGCACTGCACCTTCACCCTGGAGCGTGTCAGCCCCAGCACTAGTGACCTGGCC 1254

Qy     2284 TGCAAGCTGTGGGTGTGGCAGGTGGAGGGCGACGGGCAGAGCTTCAGCATCAACTTCAAC 2343
                   ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db     1255 TGCAAGCTGTGGGTGTGGCAGGTGGAGGGCGACGGGCAGAGCTTCAGCATCAACTTCAAC 1314

Qy     2344 ATCACCAAGGACACAAGGTTTGCTGAGCTGCTGGCTCTGGAGAGTGAAGCGGGGGTCCCA 2403
                   ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
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Db      1315 ATCACCAAGGACACAAGGTTTGCTGAGCTGCTGGCTCTGGAGAGTGAAGCGGGGGTCCCA 1374

Qy      2404 GCCCTGGTGGGCCCCAGTGCCTTCAAGATCCCCTTCCTCATTCGGCAGAAGATAATTTCC 2463
             ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db      1375 GCCCTGGTGGGCCCCAGTGCCTTCAAGATCCCCTTCCTCATTCGGCAGAAGATAATTTCC 1434

Qy      2464 AGCCTGGACCCACCCTGTAGGCGGGGTGCCGACTGGCGGACTCTGGCCCAGAAACTCCAC 2523
             ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db      1435 AGCCTGGACCCACCCTGTAGGCGGGGTGCCGACTGGCGGACTCTGGCCCAGAAACTCCAC 1494

Qy      2524 CTGGACAGCCATCTCAGCTTCTTTGCCTCCAAGCCCAGCCCCACAGCCATGATCCTCAAC 2583
             ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db      1495 CTGGACAGCCATCTCAGCTTCTTTGCCTCCAAGCCCAGCCCCACAGCCATGATCCTCAAC 1554

Qy      2584 CTGTGGGAGGCGCGGCACTTCCCCAACGGCAACCTCAGCCAGCTGGCTGCAGCAGTGGCT 2643
             ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db      1555 CTGTGGGAGGCGCGGCACTTCCCCAACGGCAACCTCAGCCAGCTGGCTGCAGCAGTGGCT 1614

Qy      2644 GGACTGGGCCAGCCAGACGCTGGCCTC-TTCACAGTG-TCGGAGGCTGAGTGCTGA 2697
             ||       |||||||  ||||  ||||  || |||||||||| |||||||||||||||||
Db      1615 GGGACTGGCCAGCAGGACGGTGGCTTCTTTCACAGTGTTCGGAGGCTGAGTGCTGA 1670


RESULT 5
US-08-808-982-3
; Sequence 3, Application US/08808982
; Patent No. 5939271
;   GENERAL INFORMATION:
;      APPLICANT:  Tessier-Lavigne, Marc
;      APPLICANT:  Leonardo, E. David
;      APPLICANT:  Hink, Lindsay
;      APPLICANT:  Masu, Masayuki
;      APPLICANT:  Kazuko, Keino-Masu
;      TITLE OF INVENTION:  Netrin Receptors
;      NUMBER OF SEQUENCES:  8
;      CORRESPONDENCE ADDRESS:
;         ADDRESSEE:  SCIENCE & TECHNOLOGY LAW GROUP
;         STREET:  268 BUSH STREET, SUITE 3200
;         CITY:  SAN FRANCISCO
;         STATE:  CALIFORNIA
;         COUNTRY:  USA
;         ZIP:  94104
;      COMPUTER READABLE FORM:
;         MEDIUM TYPE:  Floppy disk
;         COMPUTER:  IBM PC compatible
;         OPERATING SYSTEM:  PC-DOS/MS-DOS
;         SOFTWARE:  PatentIn Release #1.0, Version #1.30
;      CURRENT APPLICATION DATA:
;         APPLICATION NUMBER:  US/08/808,982
;         FILING DATE:
;         CLASSIFICATION:  530
;      ATTORNEY/AGENT INFORMATION:
;         NAME:  OSMAN, RICHARD A
;         REGISTRATION NUMBER:  36,627
;         REFERENCE/DOCKET NUMBER:  UC96-217
;      TELECOMMUNICATION INFORMATION:
;         TELEPHONE:  (415) 343-4341
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Query Match              31.2%;  Score 841.4;  DB 2;  Length 2831;
Best Local Similarity    60.0%;  Pred. No. 8.4e-182;
Matches 1638;  Conservative    0;  Mismatches  961;  Indels  130;  Gaps    9;


Qy       98 ACCCAGTGCCTGGTGCCAACCCGGACCTGCTTCCCCACTTCCTGGTGGAGCCCGAGGATG 157
            || |  | ||    ||    | | |||| | |||  || |||||||||| |||| || |||||||
Db      104 ACTCCTTCCCATCAGCACCCGCGGAGCAGCTGCCTCACTTCCTGCTGGAACCAGAGGATG 163


Qy      158 TGTACATCGTCAAGAACAAGCCAGTGCTGCTTGTGTGCAAGGCCGTGCCCGCCACGCAGA 217
            ||||||||| |||||||||||||||||     |    ||| |   ||| | || ||||| ||||
Db      164 CCTACATCGTAAAGAACAAGCCAGTGGAATTGCACTGCCGAGCCTTCCCTGCCACACAGA 223


Qy      218 TCTTCTTCAAGTGCAACGGGGAGTGGGTGCGCCAGGTGGACCACGTGATCGAGCGCAGCA 277
            ||| ||||||||| || || |||||||| |||||     | |||||| |   ||   |||
Db      224 TCTACTTCAAGTGTAATGGCGAGTGGGTTAGCCAGAAAGGCCACGTCACGCAGGAGAGCC 283


Qy      278 CAGACGGGAGCAGTGGGCTGCCCACCATGGAGGTCCGCATTAATGTCTCAAGGCAGCAGG 337
            || | |  || ||  |||  |       |||||  |  ||  | || || |||||||||
Db      284 TGGATGAGGCCACAGGCTTGCGAATACGAGAGGTGCAGATAGAGGTGTCGCGGCAGCAGG 343


Qy      338 TCGAGAAGGTGTTCGGGCTGGAGGAATACTGGTGCCAGTGCGTGGCATGGAGCTCCTCGG 397
            | |||  |   | || ||||| ||||| |||||||||| ||||||||||| |||||||| ||||
Db      344 TGGAGGAACTTTTTGGGCTCGAGGACTACTGGTGTCAGTGCGTGGCCTGGAGCTCTTCGG 403


Qy      398 GCACCACCAAGAGTCAGAAGGCCTACATCCGCATAGCCAGATTGCGCAAGAACTTCGAGC 457
            | |||||||||||     |||||||||||||| |||      ||||||||||||| || |
Db      404 GAACCACCAAGAGTCGCCGAGCCTACATCCGCATTGCCTACTTGCGCAAGAACTTTGACC 463


Qy      458 AGGAGCCGCTGGCCAAGGAGGTGTCCCTGGAGCAGGGCATCGTGCTGCCCTGCCGTCCAC 517
            ||||||| ||||| ||||||||  | |||| || |   ||   | |||| ||||| ||||
Db      464 AGGAGCCTCTGGCGAAGGAGGTACCCTTGGATCATGAGGTCCTTCTGCAGTGCCGCCCAC 523


Qy      518 CGGAGGGCATCCCTCCAGCCGAGGTGGAGTGGCTCCGGAACGAGGACCTGGTGGACCCGT 577
            | |||||  | |||   || |||||||| ||||||  ||| || || | | || ||||
Db      524 CAGAGGGAGTGCCTGTGGCTGAGGTGGAATGGCTCAAGAATGAAGATGTCATCGATCCCG 583


Qy      578 CCCTGGACCCCAATGTATACATCACGCGGGAGCACAGCCTGGTGGTGCGACAGGCCCGCC 637
            | | |||| | ||  |    |||| | ||  || |||| ||| | | || ||||| ||||
Db      584 CTCAGGACACTAACTTCCTGCTCACCATTGACCACAACCTCATCATCCGCCAGGCGCGCC 643


Qy      638 TTGCTGACACGGCCAACTACACCTGCGTGGCCAAGAACATCGTGGCACGTCGCCGCAGCG 697
            | | ||||| ||||||||||||| |||| ||||| || ||| |||| ||     |||||| |||
Db      644 TCTCAGACACAGCCAACTACACCTGTGTGGCAAAGAATATTGTGGCCAAGCGCCGGAGCA 703


Qy      698 CCTCCGCTGCTGTCATCGTCTACGTGAACGGTGGGTGGTCGACGTGGACCGAGTGGTCCG 757
            | |  || | ||||||||||||| |||||||| || ||||| |   ||| | || |||||
Db      704 CGACGGCCACAGTCATCGTCTATGTGAACGGAGGTTGGTCCAGCTGGGCAGAATGGTCAC 763
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Qy    758 TCTGCAGCGCCAGCTGTGGGCGCGGCTGGCAGAAACGGAGCCGGAGCTGCACCAACCCGG 817
          ||||    |  |||| || || || ||||||||||| |    |||  ||||||||||||| |
Db    764 CCTGCTCTAACCGCTGCGGCCGAGGTTGGCAGAAACGTACTAGGACCTGCACCAACCCAG 823

Qy    818 CGCCTCTCAACGGGGGCGCTTTCTGTGAGGGGCAGAATGTCCAGAAAACAGCCTGCGCCA 877
          | ||  |||||| || || || ||||| ||||| |||  |  |||||| || || ||| |||
Db    824 CCCCACTCAATGGAGGTGCCTTCTGCGAGGGACAGGCTTGCCAGAAGACGGCTTGCACCA 883

Qy    878 CCCTGTGCCCAGTAGACGGCAGCTGGAGCCCGTGGAGCAAGTGGTCGGCCTGTGGGCTGG 937
          || |||||||||| || ||     ||||     ||||||||||||||| |||||  |    |
Db    884 CCGTGTGCCCAGTGGATGGAGCGTGGACTGAGTGGAGCAAGTGGTCCGCCTGCAGCACAG 943

Qy    938 ACTGCACCCACTGGCGGAGCCGTGAGTGCTCTGACCCAGCACCCCGCAACGGAGGGGAGG 997
          | ||  | ||||||| |||||| ||||| ||||||  |  || | ||||  ||||||||| |
Db    944 AGTGTGCGCACTGGCGCAGCCGCGAGTGCATGGCACCGCCGCCCCAGAACGGAGGCCGTG 1003

Qy    998 AGTGCCAGGGCACTGACCTGGACACCCGCAACTGTACCAGTGACCTCTGTGTACACAGTG 1057
          | |||   || ||   || ||| || |  ||||| ||| ||  || || || || |  | |
Db   1004 ACTGCAGCGGGACGCTACTTGACTCCAAGAACTGCACCGATGGGCTGTGCGTGCTGAATC 1063

Qy   1058 CTTCTGGCC-----------------------------------CTGAGGACGTGGCCCTCT 1084
           |                                         | |  ||||||||| || |
Db   1064 AGAGAACTCTAAACGACCCTAAAAGCCGCCCCCTGGAGCCGTCGGGAGACGTGGCGCTGT 1123

Qy   1085 ATGTGGGCCTC---ATCGCCGTGGCCGTCTGCCTGGTCCTGCTGCTGCTTGTCCTCATCC 1141
          ||| |||||||    | ||||||  || |||| |  || || || || ||   |
Db   1124 ATGCGGGCCTCGTGGTGGCCGTCTTTGTGGTTCTGGCAGTTCTCATGGCTGTAGGAGTGA 1183

Qy   1142 TCGTTTATTGCCGGAAGAAGGAGGGGCTGGACTCAGATGTGGCTGACTCGTCCATT---C 1198
          |||| ||  |   | ||    || | ||| | || |  ||||||| ||   | |   |
Db   1184 TCGTGTACCGGAGAAACTGCCGGGACTTCGACACGGACATCACTGACTCCTCTGCTGCCC 1243

Qy   1199 TCACCTCAGGCTTCCAGCCCGTCAGCATCAAGCCCAGCAAAGCAGACAACCCCCATCTGC 1258
          ||||     ||  | ||||| ||||||| | ||||| |     |   |   ||||||| || || |
Db   1244 TCACTGGTGGTTTCCACCCCGTCAACTTCAAGACTGCAAGGCCCAGCAACCCACAGCTCC 1303

Qy   1259 T------CACCATCCAGCCGGACCTCAGCACCACCACCACCACCTACCAGGGCAGTCTCT 1312
          |       ||  ||  ||||||||| |   |||    |  || |||||| ||   | | |
Db   1304 TGCACCCATCCGCCCCTCCGGACCTAACGGCCAGTGCTGGCATCTACCGCGGACCTGTGT 1363

Qy   1313 GTCCCCGGCAGGA----------------------------------------TG 1327
           | ||| ||||||
Db   1364 ATGCCCTGCAGGACTCTGCCGACAAGATCCCTATGACTAATTCACCCCTTCTGGATCCCT 1423

Qy   1328 GGCCCAGCCCCAAGTTCCAGCTCACCAATGGGCACCTGCTCAGCCC---------CCTGG 1378
          |||||||||| |||| || || || ||       |       || || |           |||||
Db   1424 TGCCCAGCCTCAAGATCAAGGTCTATGACTCCAGCACCATCGGCTCTGGGGCTGGCCTGG 1483

Qy   1379 GTGGCGGCCGCCACACACTGCACCACAGCTCTCCCACCTCTGAGGCCGAGGAGTTCGTCT 1438
          || || | ||  |||   |        |   | ||| |  |   |  |  || ||| |
Db   1484 CTGATGGAGCCGACCTGCTGGGTGTCTTACCACCCGGTACATACCCAGGCGATTTCTCCC 1543

Qy   1439 CCCGCCTCTCCACCCAGAAC---------------------------TACTTCCGCTCCC 1471
           |  |  |  || | ||                           || ||| ||
Db   1544 GGGACACCCACTTCCTGCACCTGCGCAGCGCCAGCCTTGGTTCCCAGCACCTCCTGGGCC 1603
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Qy    1472 TGCCCCGAGGCACCAGCAACATGACCTATGGGACCTTCAACTTCCTCGGGGGCCGGCTGA 1531
           I II IIIII I IIIIII II    I  III IIIII   I III II II  IIIIII
Db    1604 TCCCTCGAGACCCCAGCAGCAGTGTCAGTGGCACCTTTGGTTGCCTGGGTGGGAGGCTGA 1663

Qy    1532 TGATCCCTAATACAGGTATCAGCCTCCTCATCCCCCCAGATGCCATACCCCGAGGGAAGA 1591
            II II    IIIII IIIIIII I  I II   I IIIIII IIII II III
Db    1664 CCATTCCCGGCACAGGGGTCAGCCTGTTGGTACCAAATGGAGCCATTCCCCAGGGCAAGT 1723

Qy    1592 TCTATGAGATCTACCTCACGCTGCACAAGCCGGAAGACGTGAGGTTGCCCCTAGCTGGCT 1651
           IIIIIII I II II   I IIIII I III I        I     I III I
Db    1724 TCTATGACTTGTATCTACGTATCAACAAGACTGAAAGCACCCTCCCACTTTCGGAAGGTT 1783

Qy    1652 GTCAGACCCTGCTGAGTCCCATCGTTAGCTGTGGACCCCCTGGCGTCCTGCTCACCCGGC 1711
            IIIII I  IIII III  II I III II III I III IIII II   III I
Db    1784 CCCAGACAGTATTGAGCCCCTCGGTGACCTGCGGGCCCACGGGCCTCCTCCTGTGCCGCC 1843

Qy    1712 CAGTCATCCTGGCTATGGACCACTGTGGGGAGCCCAGCCCTGACAGCTGGAGCCTGCGCC 1771
           I II  IIII II II IIIIIIII II  II   I I   IIIII I I I  I
Db    1844 CTGTTGTCCTCACTGTGCCCCACTGTGCTGAAGTCATTGCCGGAGACTGGATCTTCCAGC 1903

Qy    1772 TCAAAAAGCAGTCGTGCGAGGGCAGCTGGGAGGATGTGCTGCACCTGGGCGAGGAGGCGC 1831
           IIII I  III I      IIIII  IIIIIIIII III II     III IIIIII I I
Db    1904 TCAAGACCCAGGCCCATCAGGGCCACTGGGAGGAGGTGGTGACTTTGGATGAGGAGACTC 1963

Qy    1832 CCTCCCACCTCTACTACTGCCAGCTGGAGGCCAGTGCCTGCTACGTCTTCACCGAGCAGC 1891
            I  II II IIIIIIIIIIIII IIIII I   IIIII II II I   II IIII
Db    1964 TGAACACCCCCTGCTACTGCCAGCTAGAGGCTAAATCCTGCCACATCCTGTTGGACCAGC 2023

Qy    1892 TGGGCCGCTTTGCCCTGGTGGGAGAGGCCCTCAGCGTGGCTGCCGCCAAGCGCCTCAAGC 1951
           IIII   II I  I    III III II I I    I II IIIIII III III
Db    2024 TGGGTACCTACGTGTTCACGGGCGAGTCCTACTCCCGCTCCGCAGTCAAGCGGCTCCAGC 2083

Qy    1952 TGCTTCTGTTTGCGCCGGTGGCCTGCACCTCCCTCGAGTACAACATCCGGGTCTACTGCC 2011
           I      I II II II I   IIIIIIIIIIII IIIIII I   II IIIIIIIIII I
Db    2084 TAGCCATCTTCGCCCCAGCCCTCTGCACCTCCCTGGAGTATAGTCTCAGGGTCTACTGTC 2143

Qy    2012 TGCATGACACCCACGATGCACTCAAGGAGGTGGTGCAGCTGGAGAAGCAGCTGGGGGGAC 2071
           II I IIIIII I  I   IIIII IIIIIIII I IIIIIIIII I    IIIII II
Db    2144 TGGAGGACACTCCTGCAGCACTGAAGGAGGTCCTAGAGCTGGAGAGGACTCTGGGTGGCT 2203

Qy    2072 AGCTGATCCAGGAGCCACGGGTCCTGCACTTCAAGGACAGTTACCACAACCTGCGCCTAT 2131
           I  II I  IIIIIII   I   III III IIIIIIIIIIIIIIIIIIIIII II II I
Db    2204 ACTTGGTGGAGGAGCCCAAGACTTTGCTCTTTAAGGACAGTTACCACAACCTACG-CTCT 2262

Qy    2132 CCATCCACGATGTGCCCAGCTCCCTGTGGAAGAGTAAGCTCCTTGTCAGCTACCAGGAGA 2191
           II IIIII II  I III    III  IIII III II II II I..-II IIIIIIIIII
Db    2263 CCCTCCATGACATCCCCATGCCCACTGGAGGAGCAAACTACTGGCCAAGTACCAGGAGA 2322

Qy    2192 TCCCCTTTTATCACATCTGGAATGGCACGCAGCGGTACTTGCACTGCACCTTCACCCTGG 2251
           I IIIIII II II I IIIII IIII  III      I IIIIIIIII IIIIIIIIII
Db    2323 TTCCCTTCTACCATGTGTGGAACGGCAGCCAGAAAGCCCTGCACTGCACTTTCACCCTGG 2382

Qy    2252 AGCGTGTCAGCCCCAGCACTAGTGACCTGGCCTGCAAGCTGTGGGTGTGGCAGGTGGAGG 2311
           II I   IIII    I I I III I IIII III I II III IIIIIIII II I
Db    2383 AGAGACATAGCCTAGCCTCCACTGAGTTCACCTGTAAGGTCTGCGTGCGGCAGGTAGAAG 2442

Qy    2312 GCGACGGGCAGAGCTTCAGCATCAACTTCAACATCAC---CAAGGACACAAGGTTTGCTG 2368
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                 |  || || ||||  |||      |  |'|  ||   |  |      ||     |  | |      |
Db      . 2443 GGGAAGGCCAGATTTTCCAGCTGCACACCACGCTGGCTGAGACGCCTGCTGGCTCCCTGG 2502

Qy        2369 AGCTGCTGGCTCTGGAGAGTGAAGCGGGGGTCCCAGCCCTGGTGGGCCCCAGTGCCTTCA 2428
                 |    ||       .|    ||     |  | | |  | | | |||| |||   ||||||||
Db        2503 ATGCACTCTGCTCTGCCCCTGGCAATGCTGCCACCACACAGCTGGGACCCTATGCCTTCA 2562

Qy        2429 AGATCCCCTTCCTCATTCGGCAGAAGATAATTTCCAGCCTGGACCCACCCTGTAGGCGGG 2488
                 |||| ||  |   ||| || |||||||    |||||||||| | |||       ||||
Db        2563 AGATACCACTGTCCATCCGCCAGAAGATCTGCAACAGCCTGGACGCCCCCAACTCACGGG 2622

Qy        2489 GTGCCGACTGGCGGACTCTGGCCCAGAAACTCCACCTGGACAGCCATCTCAGCTTCTTTG 2548
                 |     ||||||||||     |||| ||||| |||   | |||||| | |  || |  || ||| |
Db        2623 GCAATGACTGGCGGCTGTTGGCACAGAAGCTCTCCATGGACCGGTACCTGAACTACTTCG 2682

Qy        2549 CCTCCAAGCCCAGCCCCACAGCCATGATCCTCAACCTGTGGGAGGCGCGGCACTTCCCCA 2608
                 ||  ||||   | ||  |||||||| |  ||||| |   ||||  ||||| ||  |||||
Db        2683 CCACCAAAGCTAGTCCCACAGGCGTGATCTTAGACCTCTGGGAAGCTCGGCAGCAGGATG 2742

Qy        2609 ACGGCAACCTCAGCCAGCTGGCTGCAGCAGTGGCTGGACTGGGCCAGCCAGACGCTGGCC 2668
                 | ||  |||||| |    |||||     ||  |||   |    ||||| ||    ||
Db        2743 ATGGGGACCTCAACAGCCTGGCCAGTGCCTTGGAGGAGATGGGCAAGAGTGAGATGCTGG 2802

Qy        2669 TCTTCACAGTGTCGGAGGCTGAGTGCTGA 2697
                 |   ||        | || |   || ||||||
Db        2803 TAGCCATGACCACTGATGGCGATTGCTGA 2831



RESULT 6
US-09-306-902A-3
; Sequence 3, Application US/09306902A
; Patent No. 6277585
;   GENERAL INFORMATION:
;       APPLICANT: Tessier-Lavigne, Marc
;               Leonardo, E. David
;               Hink, Lindsay
;               Masu, Masayuki
;               Kazuko, Keino-Masu
;       TITLE OF INVENTION: Netrin Receptors
;       NUMBER OF SEQUENCES: 9
;       CORRESPONDENCE ADDRESS:
;           ADDRESSEE: SCIENCE & TECHNOLOGY LAW GROUP
;           STREET: 268 BUSH STREET, SUITE 3200
;           CITY: SAN FRANCISCO
;           STATE: CALIFORNIA
;           COUNTRY: USA
;           ZIP: 94104
;       COMPUTER READABLE FORM:
;           MEDIUM TYPE: Floppy disk
;           COMPUTER: IBM PC compatible
;           OPERATING SYSTEM: PC-DOS/MS-DOS
;           SOFTWARE: PatentIn Release #1.0, Version #1.30
;       CURRENT APPLICATION DATA:
;           APPLICATION NUMBER: US/09/306,902A
;           FILING DATE: 07-May-1999
;           CLASSIFICATION: <Unknown>
;       ATTORNEY/AGENT INFORMATION:
```

;   NAME: OSMAN, RICHARD A
;   REGISTRATION NUMBER: 36,627
;   REFERENCE/DOCKET NUMBER: UC96-217
;   TELECOMMUNICATION INFORMATION:
;   TELEPHONE: (415) 343-4341
;   TELEFAX: (415) 343-4342
;   INFORMATION FOR SEQ ID NO: 3:
;   SEQUENCE CHARACTERISTICS:
;   LENGTH: 2831 base pairs
;   TYPE: nucleic acid
;   STRANDEDNESS: double
;   TOPOLOGY: linear
;   MOLECULE TYPE: cDNA
;   SEQUENCE DESCRIPTION: SEQ ID NO: 3:
US-09-306-902A-3

     Query Match   31.2%;   Score 841.4;   DB 3;   Length 2831;
     Best Local Similarity   60.0%;   Pred. No. 8.4e-182;
     Matches 1638; Conservative   0; Mismatches 961; Indels 130; Gaps   9;

Qy   98 ACCCAGTGCCTGGTGCCAACCCGGACCTGCTTCCCCACTTCCTGGTGGAGCCCGAGGATG 157
             || | | || || | |||| | ||| || |||||||||| |||| || |||||||
Db   104 ACTCCTTCCCATCAGCACCCGCGGAGCAGCTGCCTCACTTCCTGCTGGAACCAGAGGATG 163

Qy   158 TGTACATCGTCAAGAACAAGCCAGTGCTGCTTGTGTGCAAGGCCGTGCCCGCCACGCAGA 217
             |||||||| ||||||||||||||| | ||| ||| | || |||.|| ||||
Db   164 CCTACATCGTAAAGAACAAGCCAGTGGAATTGCACTGCCGAGCCTTCCCTGCCACACAGA 223

Qy   218 TCTTCTTCAAGTGCAACGGGGAGTGGGTGCGCCAGGTGGACCACGTGATCGAGCGCAGCA 277
             ||| ||||||||| || || ||||||||| ||||| | ||||||| | |||
Db   224 TCTACTTCAAGTGTAATGGCGAGTGGGTTAGCCAGAAAGGCCACGTCACGCAGGAGAGCC 283

Qy   278 CAGACGGGAGCAGTGGGCTGCCCACCATGGAGGTCCGCATTAATGTCTCAAGGCAGCAGG 337
             || | | || || ||| | | ||||| | || | || | || || |||||||||
Db   284 TGGATGAGGCCACAGGCTTGCGAATACGAGAGGTGCAGATAGAGGTGTCGCGGCAGCAGG 343

Qy   338 TCGAGAAGGTGTTCGGGCTGGAGGAATACTGGTGCCAGTGCGTGGCATGGAGCTCCTCGG 397
             | ||| | | || ||||| ||||| ||||||| |||||||||| |||||||||| ||||
Db   344 TGGAGGAACTTTTTGGGCTCGAGGACTACTGGTGTCAGTGCGTGGCCTGGAGCTCTTCGG 403

Qy   398 GCACCACCAAGAGTCAGAAGGCCTACATCCGCATAGCCAGATTGCGCAAGAACTTCGAGC 457
             | ||||||||||||| ||||||||||||| ||| ||||||||||||| || |
Db   404 GAACCACCAAGAGTCGCCGAGCCTACATCCGCATTGCCTACTTGCGCAAGAACTTTGACC 463

Qy   458 AGGAGCCGCTGGCCAAGGAGGTGTCCCTGGAGCAGGGCATCGTGCTGCCCTGCCGTCCAC 517
             |||||||| ||||| |||||||| || |||| || | | ||| | | |||| ||||
Db   464 AGGAGCCTCTGGCGAAGGAGGTACCCTTGGATCATGAGGTCCTTCTGCAGTGCCGCCCAC 523

Qy   518 CGGAGGGCATCCCTCCAGCCGAGGTGGAGTGGCTCCGGAACGAGGACCTGGTGGACCCGT 577
             | |||||| | ||| || |||||||||| |||||| ||| || || | | ||| ||
Db   524 CAGAGGGAGTGCCTGTGGCTGAGGTGGAATGGCTCAAGAATGAAGATGTCATCGATCCCG 583

Qy   578 CCCTGGACCCCAATGTATACATCACGCGGGAGCACAGCCTGGTGGTGCGACAGGCCCGCC 637
             | | |||| | || | |||| || |||| ||| | | | || ||||| ||||
Db   584 CTCAGGACACTAACTTCCTGCTCACCATTGACCACAACCTCATCATCCGCCAGGCGCGCC 643

Qy   638 TTGCTGACACGGCCAACTACACCTGCGTGGCCAAGAACATCGTGGCACGTCGCCGCAGCG 697

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                     |   |  ||||||  |||||||||||||||||  ||||| |||||| || |||||         ||||| |||
Db     644 TCTCAGACACAGCCAACTACACCTGTGTGGCAAAGAATATTGTGGCCAAGCGCCGGAGCA 703

Qy     698 CCTCCGCTGCTGTCATCGTCTACGTGAACGGTGGGTGGTCGACGTGGACCGAGTGGTCCG 757
                     |   |  ||   |  |||||||||||  ||||||||| || ||||| |  ||| | || |||||
Db     704 CGACGGCCACAGTCATCGTCTATGTGAACGGAGGTTGGTCCAGCTGGGCAGAATGGTCAC 763

Qy     758 TCTGCAGCGCCAGCTGTGGGCGCGGCTGGCAGAAACGGAGCCGGAGCTGCACCAACCCGG 817
                     ||||         |  ||||  || ||  || |||||||||| |    ||| ||||||||||| |
Db     764 CCTGCTCTAACCGCTGCGGCCGAGGTTGGCAGAAACGTACTAGGACCTGCACCAACCCAG 823

Qy     818 CGCCTCTCAACGGGGGCGCTTTCTGTGAGGGGCAGAATGTCCAGAAAACAGCCTGCGCCA 877
                     | || |||||| || || || ||||| ||||| |||  |  |||||| || || ||| |||
Db     824 CCCCACTCAATGGAGGTGCCTTCTGCGAGGGACAGGCTTGCCAGAAGACGGCTTGCACCA 883

Qy     878 CCCTGTGCCCAGTAGACGGCAGCTGGAGCCCGTGGAGCAAGTGGTCGGCCTGTGGGCTGG 937
                     || |||||||||| || ||     ||||     ||||||||||||||||||| |||||  |    |
Db     884 CCGTGTGCCCAGTGGATGGAGCGTGGACTGAGTGGAGCAAGTGGTCCGCCTGCAGCACAG 943

Qy     938 ACTGCACCCACTGGCGGAGCCGTGAGTGCTCTGACCCAGCACCCCGCAACGGAGGGGAGG 997
                     | ||  |  ||||||||| |||||  ||||||   |   || |  ||||  |||||||| |
Db     944 AGTGTGCGCACTGGCGCAGCCGCGAGTGCATGGCACCGCCGCCCCAGAACGGAGGCCGTG 1003

Qy     998 AGTGCCAGGGCACTGACCTGGACACCCGCAACTGTACCAGTGACCTCTGTGTACACAGTG 1057
                     | ||||   || ||   || |||  ||  ||||| |||  || || || || || | | | |
Db    1004 ACTGCAGCGGGACGCTACTTGACTCCAAGAACTGCACCGATGGGCTGTGCGTGCTGAATC 1063

Qy    1058 CTTCTGGCC----------------------------------CTGAGGACGTGGCCCTCT 1084
                     |                                          | |  |||||||| || |
Db    1064 AGAGAACTCTAAACGACCCTAAAAGCCGCCCCCTGGAGCCGTCGGGAGACGTGGCGCTGT 1123

Qy    1085 ATGTGGGCCTC---ATCGCCGTGGCCGTCTGCCTGGTCCTGCTGCTGCTTGTCCTCATCC 1141
                     |||  |||||||   |  |||||    ||   |||   | || || ||| |
Db    1124 ATGCGGGCCTCGTGGTGGCCGTCTTTGTGGTTCTGGCAGTTCTCATGGCTGTAGGAGTGA 1183

Qy    1142 TCGTTTATTGCCGGAAGAAGGAGGGGCTGGACTCAGATGTGGCTGACTCGTCCATT---C 1198
                     ||||  || |  | | ||        ||   | ||| | ||  |  |||||||  ||   | |
Db    1184 TCGTGTACCGGAGAAACTGCCGGGACTTCGACACGGACATCACTGACTCCTCTGCTGCCC 1243

Qy    1199 TCACCTCAGGCTTCCAGCCCGTCAGCATCAAGCCCAGCAAAGCAGACAACCCCCATCTGC 1258
                     ||||      || ||||| ||||||| | |||||.|   |   |   |||||| || || |
Db    1244 TCACTGGTGGTTTCCACCCCGTCAACTTCAAGACTGCAAGGCCCAGCAACCCACAGCTCC 1303

Qy    1259 T------CACCATCCAGCCGGACCTCAGCACCACCACCACCACCTACCAGGGCAGTCTCT 1312
                     |          ||  ||  |||||||||| |   ||| |   || ||||| ||    ||| |
Db    1304 TGCACCCATCCGCCCCTCCGGACCTAACGGCCAGTGCTGGCATCTACCGCGGACCTGTGT 1363

Qy    1313 GTCCCCGGCAGGA-----------------------------------------------TG 1327
                     |  |,|| ||||||
Db    1364 ATGCCCTGCAGGACTCTGCCGACAAGATCCCTATGACTAATTCACCCCTTCTGGATCCCT 1423

Qy    1328 GGCCCAGCCCCAAGTTCCAGCTCACCAATGGGCACCTGCTCAGCCC---------CCTGG 1378
                     |||||||| |||| || || ||  |      |       || || |         ||||||
Db    1424 TGCCCAGCCTCAAGATCAAGGTCTATGACTCCAGCACCATCGGCTCTGGGGCTGGCCTGG 1483

Qy    1379 GTGGCGGCCGCCACACACTGCACCACAGCTCTCCCACCTCTGAGGCCGAGGAGTTCGTCT 1438
                     ||  ||   | ||    |||      |      | ||  | |  | |  || ||| |
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Db    1484 CTGATGGAGCCGACCTGCTGGGTGTCTTACCACCCGGTACATACCCAGGCGATTTCTCCC 1543

Qy    1439 CCCGCCTCTCCACCCAGAAC-----------------------------TACTTCCGCTCCC 1471
                |  |  |  || | ||                              || |||   ||
Db    1544 GGGACACCCACTTCCTGCACCTGCGCAGCGCCAGCCTTGGTTCCCAGCACCTCCTGGGCC 1603

Qy    1472 TGCCCCGAGGCACCAGCAACATGACCTATGGGACCTTCAACTTCCTCGGGGGCCGGCTGA 1531
                | || |||| | ||||||| ||    |   ||| |||||    | ||| || ||  |||||
Db    1604 TCCCTCGAGACCCCAGCAGCAGTGTCAGTGGCACCTTTGGTTGCCTGGGTGGGAGGCTGA 1663

Qy    1532 TGATCCCTAATACAGGTATCAGCCTCCTCATCCCCCCAGATGCCATACCCGAGGGAAGA 1591
              '  || ||      ||||| ||||||| |   | ||    |  ||||| ||||  || |||
Db    1664 CCATTCCCGGCACAGGGGTCAGCCTGTTGGTACCAAATGGAGCCATTCCCCAGGGCAAGT 1723

Qy    1592 TCTATGAGATCTACCTCACGCTGCACAAGCCGGAAGACGTGAGGTTGCCCCTAGCTGGCT 1651
              |||||||  | || ||    |   ||||| | ||| |        |      |   || |
Db    1724 TCTATGACTTGTATCTACGTATCAACAAGACTGAAAGCACCCTCCCACTTTCGGAAGGTT 1783

Qy    1652 GTCAGACCCTGCTGAGTCCCATCGTTAGCTGTGGACCCCCTGGCGTCCTGCTCACCCGGC 1711
              |||||   |   |||| |||   || | ||| || ||| | ||| |||| ||    ||| |
Db    1784 CCCAGACAGTATTGAGCCCCTCGGTGACCTGCGGGCCCACGGGCCTCCTCCTGTGCCGCC 1843

Qy    1712 CAGTCATCCTGGCTATGGACCACTGTGGGGAGCCCAGCCCTGACAGCTGGAGCCTGCGCC 1771
              | ||  |||| || ||  |   ||||||| ||    || ||  || ||||| | | |  |
Db    1844 CTGTTGTCCTCACTGTGCCCCACTGTGCTGAAGTCATTGCCGGAGACTGGATCTTCCAGC 1903

Qy    1772 TCAAAAAGCAGTCGTGCGAGGGCAGCTGGGAGGATGTGCTGCACCTGGGCGAGGAGGCGC 1831
              ||||  |   ||| |       |||||  |||||||||  ||| || |||||| | |
Db    1904 TCAAGACCCAGGCCCATCAGGGCCACTGGGAGGAGGTGGTGACTTTGGATGAGGAGACTC 1963

Qy    1832 CCTCCCACCTCTACTACTGCCAGCTGGAGGCCAGTGCCTGCTACGTCTTCACCGAGCAGC 1891
                |   || || ||||||||||||||| |||||  |     |||||  || || |    || ||||
Db    1964 TGAACACCCCCTGCTACTGCCAGCTAGAGGCTAAATCCTGCCACATCCTGTTGGACCAGC 2023

Qy    1892 TGGGCCGCTTTGCCCTGGTGGGAGAGGCCCTCAGCGTGGCTGCCGCCAAGCGCCTCAAGC 1951
              ||||      ||  |    |    ||| ||| || |  |     | || | |||||| ||| |||
Db    2024 TGGGTACCTACGTGTTCACGGGCGAGTCCTACTCCCGCTCCGCAGTCAAGCGGCTCCAGC 2083

Qy    1952 TGCTTCTGTTTGCGCCGGTGGCCTGCACCTCCCTCGAGTACAACATCCGGGTCTACTGCC 2011
              |      | || || || | |       ||||||||||||| |||||  |  || ||||||||| |
Db    2084 TAGCCATCTTCGCCCCAGCCCTCTGCACCTCCCTGGAGTATAGTCTCAGGGTCTACTGTC 2143

Qy    2012 TGCATGACACCCACGATGCACTCAAGGAGGTGGTGCAGCTGGAGAAGCAGCTGGGGGGAC 2071
              || | ||||| |  |     ||||| ||||||||| |  ||||||||| |     ||||| ||
Db    2144 TGGAGGACACTCCTGCAGCACTGAAGGAGGTCCTAGAGCTGGAGAGGACTCTGGGTGGCT 2203

Qy    2072 AGCTGATCCAGGAGCCACGGGTCCTGCACTTCAAGGACAGTTACCACAACCTGCGCCTAT 2131
              |   || |  |||||||      |   ||| ||| ||||||||||||||||||||||| || || |
Db    2204 ACTTGGTGGAGGAGCCCAAGACTTTGCTCTTTAAGGACAGTTACCACAACCTACG-CTCT 2262

Qy    2132 CCATCCACGATGTGCCCAGCTCCCTGTGGAAGAGTAAGCTCCTTGTCAGCTACCAGGAGA 2191
              || ||||| ||   | |||      ||| |||| ||| ||  |  || | |  |||||||||||
Db    2263 CCCTCCATGACATCCCCCATGCCCACTGGAGGAGCAAACTACTGGCCAAGTACCAGGAGA 2322

Qy    2192 TCCCCTTTTATCACATCTGGAATGGCACGCAGCGGTACTTGCACTGCACCTTCACCCTGG 2251
              | |||||| || || | ||||| |||| |||   |      | ||||||||||| |||||||||||
Db    2323 TTCCCTTCTACCATGTGTGGAACGGCAGCCAGAAAGCCCTGCACTGCACTTTCACCCTGG 2382
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Qy      2252 AGCGTGTCAGCCCCAGCACTAGTGACCTGGCCTGCAAGCTGTGGGTGTGGCAGGTGGAGG 2311
             || |    ||||    | | | ||| |  |||| ||| | || ||| ||||||| || |
Db      2383 AGAGACATAGCCTAGCCTCCACTGAGTTCACCTGTAAGGTCTGCGTGCGGCAGGTAGAAG 2442

Qy      2312 GCGACGGGCAGAGCTTCAGCATCAACTTCAACATCAC---CAAGGACACAAGGTTTGCTG 2368
             | || || |||| |||  | || || | |  | ||  |  || |  | |
Db      2443 GGGAAGGCCAGATTTTCCAGCTGCACACCACGCTGGCTGAGACGCCTGCTGGCTCCCTGG 2502

Qy      2369 AGCTGCTGGCTCTGGAGAGTGAAGCGGGGGTCCCAGCCCTGGTGGGCCCCAGTGCCTTCA 2428
             |   ||     |    || | | | |  | || |||| |||||||||
Db      2503 ATGCACTCTGCTCTGCCCCTGGCAATGCTGCCACCACACAGCTGGGACCCTATGCCTTCA 2562

Qy      2429 AGATCCCCTTCCTCATTCGGCAGAAGATAATTTCCAGCCTGGACCCACCCTGTAGGCGGG 2488
             |||| ||  |   ||| || |||||||||         |||||||||| | |||     ||||
Db      2563 AGATACCACTGTCCATCCGCCAGAAGATCTGCAACAGCCTGGACGCCCCAACTCACGGG 2622

Qy      2489 GTGCCGACTGGCGGACTCTGGCCCAGAAACTCCACCTGGACAGCCATCTCAGCTTCTTTG 2548
             |  |||||||||    |||| ||||| |||  | |||||| | | || | || ||| |
Db      2623 GCAATGACTGGCGGCTGTTGGCACAGAAGCTCTCCATGGACCGGTACCTGAACTACTTCG 2682

Qy      2549 CCTCCAAGCCCAGCCCCACAGCCATGATCCTCAACCTGTGGGAGGCGCGGCACTTCCCCA 2608
             || ||||   | || ||||||| | |||||  |||| |  |||| ||||| || |||||
Db      2683 CCACCAAAGCTAGTCCCACAGGCGTGATCTTAGACCTCTGGGAAGCTCGGCAGCAGGATG 2742

Qy      2609 ACGGCAACCTCAGCCAGCTGGCTGCAGCAGTGGCTGGACTGGGCCAGCCAGACGCTGGCC 2668
             | ||  |||||| |   ||||| ||  ||| |  ||||| ||  ||
Db      2743 ATGGGGACCTCAACAGCCTGGCCAGTGCCTTGGAGGAGATGGGCAAGAGTGAGATGCTGG 2802

Qy      2669 TCTTCACAGTGTCGGAGGCTGAGTGCTGA 2697
             |   ||     | || | || ||||||
Db      2803 TAGCCATGACCACTGATGGCGATTGCTGA 2831


RESULT 7
US-09-949-016-4794
; Sequence 4794, Application US/09949016
; Patent No. 6812339
; GENERAL INFORMATION:
;   APPLICANT: VENTER, J. Craig et al.
;   TITLE OF INVENTION: POLYMORPHISMS IN KNOWN GENES ASSOCIATED
;   TITLE OF INVENTION:  WITH HUMAN DISEASE, METHODS OF DETECTION AND USES
THEREOF
;   FILE REFERENCE: CL001307
;   CURRENT APPLICATION NUMBER: US/09/949,016
;   CURRENT FILING DATE:  2000-04-14
;   PRIOR APPLICATION NUMBER: 60/241,755
;   PRIOR FILING DATE: 2000-10-20
;   PRIOR APPLICATION NUMBER: 60/237,768
;   PRIOR FILING DATE: 2000-10-03
;   PRIOR APPLICATION NUMBER: 60/231,498
;   PRIOR FILING DATE: 2000-09-08
;   NUMBER OF SEQ ID NOS: 207012
;   SOFTWARE: FastSEQ for Windows Version 4.0
; SEQ ID NO 4794
;   LENGTH: 3008
;   TYPE: DNA
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Query Match              30.9%;  Score 833.6;  DB 4;  Length 3008;
Best Local Similarity    62.1%;  Pred. No. 5.1e-180;
Matches 1435;  Conservative    0;  Mismatches  814;  Indels   63;  Gaps    5;
```

```
Qy     440 TGCGCAAGAACTTCGAGCAGGAGCCGCTGGCCAAGGAGGTGTCCCTGGAGCAGGGCATCG 499
           |  || ||||  || ||||||||| ||  || |  ||||| |||||  |||| ||||   ||
Db      10 TACGGAAGACATTTGAGCAGGAACCCCTAGGAAAGGAAGTGTCTTTGGAACAGGAAGTCT 69

Qy     500 TGCTGCCCTGCCGTCCACCGGAGGGCATCCCTCCAGCCGAGGTGGAGTGGCTCCGGAACG 559
           | ||| |  || || ||||| || || ||||||    || ||||||||| |||| |   || |
Db      70 TACTCCAGTGTCGACCACCTGAAGGGATCCCAGTGGCTGAGGTGGAATGGTTGAAAAATG 129

Qy     560 AGGACCTGGTGGACCCGTCCCTGGACCCCAATGTATACATCACGCGGGAGCACAGCCTGG 619
           | |||| | | || ||    || |  ||| | ||| | ||  || || ||  |||| |||
Db     130 AAGACATAATTGATCCCGTTGAAGATCGGAATTTTTATATTACTATTGATCACAACCTCA 189

Qy     620 TGGTGCGACAGGCCCGCCTTGCTGACACGGCCAACTACACCTGCGTGGCCAAGAACATCG 679
           |  |     ||||||||| ||  ||||  || || || || || ||||||||| || |||||| |
Db     190 TCATAAAGCAGGCCCGACTCTCTGATACTGCAAATTACACCTGTGTTGCCAAAAACATTG 249

Qy     680 TGGCACGTCGCCGCAGCGCCTCCGCTGCTGTCATCGTCTACGTGAACGGTGGGTGGTCGA 739
           |  ||      |   ||  | | | ||  |||||||| ||||| || |||||||| ||||| |
Db     250 TTGCCAAGAGGAAAAGTACAACTGCCACTGTCATAGTCTATGTCAACGGTGGCTGGTCCA 309

Qy     740 CGTGGACCGAGTGGTCCGTCTGCAGCGCCAGCTGTGGGCGCGGCTGGCAGAAACGGAGCC 799
           | ||||| |||||||| || || |  |  |||||||| || || |  |||||||| |
Db     310 CCTGGACGGAGTGGTCTGTGTGTAACAGCCGCTGTGGACGAGGGTATCAGAAACGTACAA 369

Qy     800 GGAGCTGCACCAACCCGGCGCCTCTCAACGGGGGCGCTTTCTGTGAGGGGCAGAATGTCC 859
           |||    ||  |||||||||||||| ||  |||||  ||||| ||  |||||||| ||||||| |
Db     370 GGACTTGTACCAACCCGGCACCACTCAATGGGGGTGCCTTCTGTGAAGGGCAGAGTGTGC 429

Qy     860 AGAAAACAGCCTGCGCCACCCTGTGCCCAGTAGACGGCAGCTGGAGCCCGTGGAGCAAGT 919
           |||||| |||||| || | | |||||||| || ||||| |  ||||||| | |||||||||||
Db     430 AGAAAATAGCCTGTACTACGTTATGCCCAGTGGATGGCAGGTGGACGCCATGGAGCAAGT 489

Qy     920 GGTCGGCCTGTGGGCTGGACTGCACCCACTGGCGGAGCCGTGAGTGCTCTGACCCAGCAC 979
           |||| |·|||||   || |||||||||||||||| ||  | ||||||| | |  ||||| |
Db     490 GGTCTACTTGTGGAACTGAGTGCACCCACTGGCGCAGGAGGGAGTGCACGGCGCCAGCCC 549

Qy     980 CCCGCAACGGAGGGGAGGAGTGCCAGGGCACTGACCTGGACACCCGCAACTGTACCAGTG 1039
           ||    ||  || |||||  |||| ||| ||| | |||  |  || | ||||| || || ||
Db     550 CCAAGAATGGAGGCAAGGACTGCGACGGCCTCGTCTTGCAATCCAAGAACTGCACTGATG 609

Qy    1040 ACCTCTGTGTACACAGTGCTTCTGGCCCTGAGGACGTGGCCCTCTATGTGGG---CCTCA 1096
           || || |   | || | |||| | |||   || | ||  || || || ||||||||| |  | |
Db     610 GGCTTTGCATGCAGACTGCTCCTGATTCAGATGATGTTGCTCTCTATGTTGGGATTGTGA 669

Qy    1097 TCGCCGTGGCCGTCTGCCTGGTCCTGCTGCTGCTTGTCCTCATCCTCGTTTATTGCCGGA 1156
           | || ||| ||  ||| ||||||| |   |   || |  | || ||| || |||| |  ||
Db     670 TAGCAGTGATCGTTTGCCTGGCGATCTCTGTAGTTGTGGCCTTGTTTGTGTATCGGAAGA 729

Qy    1157 AGAAGGAGGGGCTGGACTCAGATGTGGCTGACTCGTCCATTCTCACCTCAGGCTTCCAGC 1216
           | |     |   |  || ||||||| |   |||||| ||     ||||      ||||| ||||
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```
Db    730 ATCATCGTGACTTTGAGTCAGATATTATTGACTCTTCGGCACTCAATGGGGGCTTTCAGC 789

Qy   1217 CCGTCAGCATCAAGCCCAGCAAAGCAGACAACCCCCATCTGCTCACCATCCAGCCGGACC 1276
           | || |  |||||||        |||| |  |    ||||||| |  |  |  || ||||
Db    790 CTGTGAACATCAAG--------GCAGCAAGACAAGATCTGCTGGCTGTACCCCCAGACC  840

Qy   1277 TCAGCACCACCACCACCACCTACCAGGGCAGTCTCTGTCCCCGGCAGGATG-----GGCC 1331
           |||    |  |  |   |||  |||  ||     ||  | |||  | ||| ||  ||   | |
Db    841 TCACGTCAGCTGCAGCCATGTACAGAGGACCTGTCTATGCCCTGCATGACGTCTCAGACA  900

Qy   1332 CAGCCCCAAGTTCCAGCTCACCAAT----GGGCACCTGCTCAGCCCCCTGGGTGGCGGCC 1387
           |  |||||  ||| ||| |||||||     |  |  |||| || || |           |
Db    901 AAATCCCAATGACCAACTCTCCAATTCTGGATCCACTGCCCAACCTGAAAATCAAAGTGT  960

Qy   1388 GCCACACACTGCACCACAGCTCTCCCACCTCTGAGGCCGAGGAGTTCGTCTCCCGCCTCT 1447
           |  ||||        |  | |||    |||   |  |||||     | ||  ||  || |
Db    961 ACAACACCTCAGGTGCTGTCACCCCCCAAGATGACCTCTCTGAGTTTACGTCCAAGCTGT 1020

Qy   1448 CCACCCAGAACTACTTCCGCTCCCTGCCCCGAGGCACCAGCAACATGA------------ 1495
           || | ||||    |     | |   ||    |   || || || | |||
Db   1021 CCCCTCAGATGACCCAGTCGTTGTTGGAGAATGAAGCCCTCAGCCTGAAGAACCAGAGTC 1080

Qy   1496 -------------------------------CCTATGGGACCTTCAACTTCCTCGGGGGCC 1525
                                           |  | |||  | |||||||||  ||  || |
Db   1081 TAGCAAGGCAGACTGATCCATCCTGTACCGCATTTGGCAGCTTCAACTCGCTGGGAGGTC 1140

Qy   1526 GGCTGATGATCCCTAATACAGGTATCAGCCTCCTCATCCCCCCAGATGCCATACCCCGAG 1585
           || ||  | || ||| ||||     ||||| |||||| || ||| | |   ||||| |||| ||
Db   1141 ACCTTATTGTTCCCAATTCAGGAGTCAGCTTGCTGATTCCCGCTGGGGCCATTCCCCAAG 1200

Qy   1586 GGAAGATCTATGAGATCTACCTCACGCTGCACAAGCCGGAAGACGTGAGGTTGCCCCTAG 1645
           |||    ||||  ||  ||  ||  | ||  | |||| |   |||      |||||   |||  | |
Db   1201 GGAGAGTCTACGAAATGTATGTGACTGTACACAGGAAAGAAACTATGAGGCCACCCATGG 1260

Qy   1646 CTGGCTGTCAGACCCTGCTGAGTCCCATCGTTAGCTGTGGACCCCCTGGCGTCCTGCTCA 1705
           || || |||||| ||  ||   || | || |||||||||| ||||| || |  |||||||
Db   1261 ATGACTCTCAGACACTTTTGACCCCTGTGGTGAGCTGTGGGCCCCCAGGAGCTCTGCTCA 1320

Qy   1706 CCCGGCCAGTCATCCTGGCTATGGACCACTGTGGGGAGCCCAGCCCTGACAGCTGGAGCC 1765
           |||| |||||| | ||| |||||| |  ||| |  | |  || | |||||     |||||
Db   1321 CCCGCCCAGTCGTCCTCACTATGCATCACTGCGCAGACCCCAATACCGAGGACTGGAAAA 1380

Qy   1766 TGCGCCTCAAAAAGCAGTCGTGCGAGGGCAGCTGGGAGGATGTGCTGCACCTGGGCGAGG 1825
           | |   ||||| || || |||  |    |||| |||||||||||| ||     | || ||||
Db   1381 TACTGCTCAAGAACCAGGCAGCACAGGGACAGTGGGAGGATGTGGTGGTGGTCGGGGAGG 1440

Qy   1826 AGGCGCCCTCCCACCTCTACTACTGCCAGCTGGAGGCCAGTGCCTGCTACGTCTTCACCG 1885
           |         | ||  || || ||||    |||||||| ||      ||||||  || || |
Db   1441 AAAACTTCACCACCCCCTGCTACATTCAGCTGGATGCAGAGGCCTGCCACATCCTCACAG 1500

Qy   1886 AGCAGCTGGGCCGCTTTGCCCTGGTGGGAGAGGCCCTCAGCGTGGCTGCCGCCAAGCGCC 1945
           || | ||  || || ||||||||| ||| |  || ||  |  || || || |||||||
Db   1501 AGAACCTCAGCACCTACGCCCTGGTAGGACATTCCACCACCAAAGCGGCTGCGAAGCGCC 1560

Qy   1946 TCAAGCTGCTTCTGTTTGCGCCGGTGGCCTGCACCTCCCTCGAGTACAACATCCGGGTCT 2005
           |||||||||    | |||| ||| || |||| |||| |||| ||||||||| |||||| ||||
Db   1561 TCAAGCTGGCCATCTTTGGGCCCCTGTGCTGCTCCTCGCTGGAGTACAGCATCCGAGTCT 1620
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Qy    2006 ACTGCCTGCATGACACCCACGATGCACTCAAGGAGGTGGTGCAGCTGGAGAAGCAGCTGG 2065
           |||| ||| |||||||||| |||||| || |||||| | | || || |||| ||| |||
Db    1621 ACTGTCTGGATGACACCCAGGATGCCCTGAAGGAAATTTTACATCTTGAGAGACAGATGG 1680

Qy    2066 GGGGACAGCTGATCCAGGAGCCACGGGTCCTGCACTTCAAGGACAGTTACCACAACCTGC 2125
           | ||||||||| | | || || || || || || || | ||| ||||||||||||
Db    1681 GAGGACAGCTCCTAGAAGAACCTAAGGCTCTTCATTTTAAAGGCAGCACCCACAACCTGC 1740

Qy    2126 GCCTATCCATCCACGATGTGCCCAGCTCCCTGTGGAAGAGTAAGCTCCTTGTCAGCTACC 2185
           |||| || || ||||||| | || ||||| |||||||| || | ||| | | || |
Db    1741 GCCTGTCAATTCACGATATCGCCCATTCCCTCTGGAAGAGCAAATTGCTGGCTAAATATC 1800

Qy    2186 AGGAGATCCCCTTTTATCACATCTGGAATGGCACGCAGCGGTACTTGCACTGCACCTTCA 2245
           |||| || || |||||| || | |||| ||| | || | || |||||||||||||||||
Db    1801 AGGAAATTCCATTTTACCATGTTTGGAGTGGATCTCAAAGAAACCTGCACTGCACCTTCA 1860

Qy    2246 CCCTGGAGCGTGTCAGCCCCAGCACTAGTGACCTGGCCTGCAAGCTGTGGGTGTGGCAGG 2305
           | ||||| | | |||| | ||| || |||| ||||| || || ||| ||||||
Db    1861 CTCTGGAAAGATTTAGCCTGAACACAGTGGAGCTGGTTTGCAAACTCTGTGTGCGGCAGG 1920

Qy    2306 TGGAGGGCGACGGGCAGAGCTTCAGCATCAACTTCAACATCACCAAGGACACAAGGTTTG 2365
           |||| || || ||||||| |||| |||||| || | | | |||| | |
Db    1921 TGGAAGGAGAAGGGCAGATCTTCCAGCTCAACTGCACCGTGTCAGAGGAACCTACTGGCA 1980

Qy    2366 CTGAGCTGCTGGCTCTGGAGAGTGAAGCGGGGGTCCCAGCCCTGGTGGGCCCCAGTGCCT 2425
           || ||| | ||||| || || | | | ||| |||||||| |
Db    1981 TCGATTTGCCGCTGCTGGATCCTGCGAACACCATCACCACGGTCACGGGGCCCAGTGCTT 2040

Qy    2426 TCAAGATCCCCTTCCTCATTCGGCAGAAGATAATTTCCAGCCTGGACCCACCCTGTAGGC 2485
           ||| ||||| ||| || ||||||||| | | ||||||||| | ||| | |
Db    2041 TCAGCATCCCTCTCCCTATCCGGCAGAAGCTCTGTAGCAGCCTGGATGCCCCCCAGACGA 2100

Qy    2486 GGGGTGCCGACTGGCGGACTCTGGCCCAGAAACTCCACCTGGACAGCCATCTCAGCTTCT 2545
           | || ||||||| ||| ||||||||| || || |||||||||| | | | | ||
Db    2101 GAGGCCATGACTGGAGGATGCTGGCCCATAAGCTGAACCTGGACAGGTACTTGAATTACT 2160

Qy    2546 TTGCCTCCAAGCCCAGCCCCACAGCCATGATCCTCAACCTGTGGGAGGCGCGGCACTTCC 2605
           ||||| |||| ||||||| || | | | |||||| | || |||||| || | | ||||||
Db    2161 TTGCCACCAAATCCAGCCCAACTGGCGTAATCCTGGATCTTTGGGAAGCACAGAACTTCC 2220

Qy    2606 CCAACGGCAACCTCAGCCAGCTGGCTGCAGCAGTGGCTGGACTGGGCCAGCCAGACGCTG 2665
           | | || ||||| ||| ||||| || | ||| | | |||| | || | |
Db    2221 CAGATGGAAACCTGAGCATGCTGGCAGCTGTCTTGGAAGAAATGGGAAGACATGAAACGG 2280

Qy    2666 GCCTCTTCACAGTGTCGGAGGCTGAGTGCTGA 2697
           | | | || | || | ||| | |
Db    2281 TGGTGTCCTTAGCAGCAGAAGGGCAGTATTAA 2312
```

RESULT 8
US-09-969-532-9
; Sequence 9, Application US/09969532
; Patent No. 6777232
; GENERAL INFORMATION:
; APPLICANT: Walke, D. Wade
; APPLICANT: Scoville, John

; TITLE OF INVENTION: No. 6777232el Human Membrane Proteins and Polynucleotides Encoding the Same
; FILE REFERENCE: LEX-0244-USA
; CURRENT APPLICATION NUMBER: US/09/969,532
; CURRENT FILING DATE: 2001-10-02
; PRIOR APPLICATION NUMBER: US 60/237,280
; PRIOR FILING DATE: 2000-10-02
; NUMBER OF SEQ ID NOS: 33
; SOFTWARE: FastSEQ for Windows Version 4.0
; SEQ ID NO 9
;   LENGTH: 2736
;   TYPE: DNA
;   ORGANISM: homo sapiens
US-09-969-532-9

    Query Match   18.1%;   Score 487; DB 4; Length 2736;
    Best Local Similarity 50.9%;   Pred. No. 4.1e-101;
    Matches 1305; Conservative   0; Mismatches 1230; Indels   30; Gaps   5;

Qy   127 CTTCCCCACTTCCTGGTGGAGCCCGAGGATGTGTACATCGTCAAGAACAAGCCAGTGCTG 186
              || || || ||| | | |||||| || |||| || || |||||| ||| || |
Db   157 CTGCCTCATTTCATAGAGGAGCCAGATGATGCTTATATTATCAAGAGCAACCCTATTGCA 216

Qy   187 CTTGTGTGCAAGGCCGTGCCCGCCACGCAGATCTTCTTCAAGTGCAACGGGGAGTGGGTG 246
              || |||||| || ||| |||| ||||||| ||||||||| ||||||||| |||||||||
Db   217 CTCAGGTGCAAAGCGAGGCCAGCCATGCAGATATTCTTCAAATGCAACGGCGAGTGGGTC 276

Qy   247 CGCCAGGTGGACCACGTGATCGAGCGCAGCACAGACGGGAGCAGTGGGCTGCCCACCATG 306
              | ||| || ||||| || | |||| |||| || || |
Db   277 CATCAGAACGAGCACGTCTCTGAAGAGACTCTGGACGAGAGCTCAGGTTTGAAGGTCCGC 336

Qy   307 GAGGTCCGCATTAATGTCTCAAGGCAGCAGGTCGAGAAGGTGTTCGGGCTGGAGGAATAC 366
              || || ||| ||||| | ||||| ||||| ||| | | |||| ||||| ||
Db   337 GAAGTGTTCATCAATGTTACTAGGCAACAGGTGGAGGACTTCCATGGGCCCGAGGACTAT 396

Qy   367 TGGTGCCAGTGCGTGGCATGGAGCTCCTCGGGCACCACCAAGAGTCAGAAGGCCTACATC 426
              |||||||||| ||||| |||||| | ||| ||| ||||||| |||||||| |
Db   397 TGGTGCCAGTGTGTGGCGTGGAGCCACCTGGGTACCTCCAAGAGCAGGAAGGCCTCTGTG 456

Qy   427 CGCATAGCCAGATTGCGCAAGAACTTCGAGCAGGAGCCGCTGGCCAAGGAGGTGTCCCTG 486
              ||||||||| || || || ||||| || || || || || | | | ||| || || |
Db   457 CGCATAGCCTATTTACGGAAAAACTTTGAACAAGACCCACAAGGAAGGGAAGTTCCCATT 516

Qy   487 GAGCAGGGCATCGTGCTGCCCTGCCGTCCACCGGAGGGCATCCCTCCAGCCGAGGTGGAG 546
              || || || |||| ||||||| ||||| ||||| |||||| | |||||||||||
Db   517 GAAGGCATGATTGTACTGCACTGCCGCCCACCAGAGGGAGTCCCTGCTGCCGAGGTGGAA 576

Qy   547 TGGCTCCGGAACGAGGACCTGGTGGACCCGTCCCTGGACCCCAATGTATACATCACGCGG 606
              ||||| || || || || | | ||| | | ||| || | | ||| || |
Db   577 TGGCTGAAAAATGAAGAGCCCATTGACTCTGAACAAGACGAGAACATTGACACCAGGGCT 636

Qy   607 GAGCACAGCCTGGTGGTGCGACAGGCCCGCCTTGCTGACACGGCCAACTACACCTGCGTG 666
              || || | |||| | | | | |||||| || || | ||| | | || |||||||||| ||
Db   637 GACCATAACCTGATCATCAGGCAGGCACGGCTCTCGGACTCAGGAAATTACACCTGCATG 696

Qy   667 GCCAAGAACATCGTGGCACGTCGCCGCAGCGCCTCCGCTGCTGTCATCGTCTACGTGAAC 726
              || |||||||||||| | | ||| | || || |||| | |||||||||| |

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Db    697  GCAGCCAACATCGTGGCTAAGAGGAGAAGCCTGTCGGCCACTGTTGTGGTCTACGTGGAT  756

Qy    727  GGTGGGTGGTCGACGTGGACCGAGTGGTCCGTCTGCAGCGCCAGCTGTGGGCGCGGCTGG  786
           ||  |  |||     |||||  |||  ||||||||||||||||||  |     ||||
Db    757  GGGAGCTGGGAAGTGTGGAGCGAATGGTCCGTCTGCAGTCCAGAGTGTGA---------A  807

Qy    787  CAGAAACGGAGCCGGAGCTGCACCAACCCGGCGCCTCTCAACGGGGGCGCTTTCTGTGAG  846
           ||    |||| ||||     |||||    ||  |  ||    || ||||||   |||||||
Db    808  CATTTGCGGATCCGGGAGTGCACAGCACCACCCCCGAGAAATGGGGGCAAATTCTGTGAA  867

Qy    847  GGGCAGAATGTCCAGAAAACAGCCTGCGCCACCCTGTGCCCAGTAGACGGCAGCTGGAGC  906
           || |  |      |       |  |||| |                |     |    |
Db    868  GGTCTAAGCCAGGAATCTGAAAACTGCACAGATGGTCTTTGCATCCTAGATAAAAAACCT  927

Qy    907  CCGTGGAGCAAGTGGTCGGCCTGTGGGCTGGACTGCACCCACTGGCGGAGCCGTGAGTGC  966
           |           |       |        |   ·  ||  |  ||               | ||
Db    928  CTTCATGAAATAAAACCCCAAAGCATTGAGAATGCCAGCGACATTGCTTTGTACTCGGGC  987

Qy    967  TCTGACCCAGCACCCCGCAACGGAGGGGAGGAGTGCCAGGGCACTGACCTGGACACCCGC  1026
           | |   |  | ||    |        ||  |  |      |  |      ||   |||   ||  ||
Db    988  TTGGGTGCTGCCGTCGTGGCCGTTGCAGTCCTGGTCATTGGTGTCACCCTTTACAGACGG  1047

Qy   1027  AACTGTACCAGTGACCTCTGTGTACACAGTGCTTCTGGCCCTGAGGACGTGGCCCTCTAT  1086
           | |   |       |     |     |     || ||  |||    |     |   |   || |
Db   1048  AGCCAGAGTGACTATGGCGTGGACGTCATTGACTCTTCTGCATTGACAGGTGGCTTCCA-  1106

Qy   1087  GTGGGCCTCATCGCCGTGGCCGTCTGCCTGGTCCTGCTGCTGCTTGTCCTCATCCTCGTT  1146
            |  | ||| |  |   | ||| |  | ||  |      |       |  | || || |
Db   1107  --GACCTTCAACTTCAAAACAGTCCGTCAAGCCAAGAATATCATGGAACTAATGATACAA  1164

Qy   1147  TATTGCCGGAAGAAGGAGGGGCTGGACTCAGATGTGGCTGACTCGTCCATTCTCACCTCA  1206
           |    ·       |   ||| |   || | ||          | |    ||| ||
Db   1165  GAAAAATCCTTTGGTAACTCCCTGCTCCTGAATTCTGCCATGCAGCCAGATCTGACAGTG  1224

Qy   1207  GGCTTCCAGCCCGTCAGCATCAAGCCCAGCAAAGCAGACAACCCCCATCTGCTCACCATC  1266
            ||        |  |  |    ||          | |          | |    |||||  |
Db   1225  AGCCGGACATACAGCGGACCCATCTGTCTGCAGGACCCTCTGGACAAGGAGCTCATGACA  1284

Qy   1267  CAGCCGGACCTCAGCACCACCACCACCACCTACCAGGGCAGTCTCTGTCCCCGGCAGGAT  1326
           || |   |||   | |  |      |   |   | |   | ||     |  | |
Db   1285  GAGTCCTCACTCTTTAACCCTTTGTCGGACATCAAAGTGAAAGTCCAGAGCTCGTTCATG  1344

Qy   1327  GGGCCCAGCCCCAAGTTCCAGCTCACCAATGGGCACCTGCTCAGCCCCCTGGGTGGCGGC  1386
           |   ||       ||  ||   ||     |  |   |||    |     | |    |
Db   1345  GTTTCCCTGGGAGTGTCTGAGAGAGCTGAGTACCACGGCAAGAATCATTCCAGGACTTTT  1404

Qy   1387  CGCCACACACTGCACCACAGCTCTCCCACCTCTGAGGCCGAGGAGTTCGTCTCCCGCCTC  1446
           | |||   |        |||||||||| |   ||     |  ||     |     ||  | ||
Db   1405  CCCCATGGAAACAACCACAGCTTTAGTACAATGCATCCCAGAAATAAAATGCCCTACATC  1464

Qy   1447  TCCACCCAGAACTACTTCCGCTCCCTGCCCCGAGGCACCAGCAACATGACCTATGGGACC  1506
            |  |  |          || || ||| | | ||        | ||||   ||| |
Db   1465  CAAAATCTGTC---------ATCACTCCCCACAAGGACAGAACTGAGGACAACTGGTGTC  1515

Qy   1507  TTCAACTTCCTCGGGGGCCGGCTGATGATCCCTAATACAGGTATCAGCCTCCTCATCCCC  1566
           ||   |    | |||||| ||  |  | || || |||||||||  | ||| | ||||||  ||
Db   1516  TTTGGCCATTTAGGGGGGCGCTTAGTAATGCCAAATACAGGGGTGAGCTTACTCATACCA  1575
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Qy   1567 CCAGATGCCATACCCCGAGGGAAGATCTATGAGATCTACCTCACGCTGCACAAGCCGGAA 1626
          |   |  |||||| ||    |  |||    |  ||||| ||   |   |    ||| |  |
Db   1576 CACGGTGCCATCCCAGAGGAGAATTCTTGGGAGATTTATATGTCCAT---CAACCAAGGT 1632

Qy   1627 GACGTGAGGTTGCCCCTAGCTGGCTGTCAGACCCTGCTGAGTCCCATCGTTAGCTGTGGA 1686
          ||     ||  | |     || |||||| | ||   || |||||||||     || | ||||||
Db   1633 GAACCCAGCCTCCAGTCAGATGGCTCTGAGGTGCTCCTGAGTCCTGAAGTCACCTGTGGT 1692

Qy   1687 CCCCCTGGCGTCCTGCTCACCCGGCCAGTCATCCTGGCTATGGACCACTGTGGGGAGCCC 1746
          || || | | |  | |||     || ||  |     || | ||  |||||||| ||   |
Db   1693 CCTCCAGACATGATCGTCACCACTCCCTTTGCATTGACCATCCCGCACTGTGCAGATGTC 1752

Qy   1747 AGCCCTGACAGCTGGAGCCTGCGCCTCAAAAAGCAGTCGTGCGAGGGCAGCTGGGAGGAT 1806
          || |||| |||| ||  | |  | || ||| |||||||| ||||||||
Db   1753 AGTTCTGAGCATTGGAATATCCATTTAAAGAAGAGGACACAGCAGGGCAAATGGGAGGAA 1812

Qy   1807 GTGCTGCACCTGGGCGAGGAGGCGCCCTCCCACCTCTACTACTGCCAGCTGGAGGCCAGT 1866
          ||| ||   ||| || || |  |   ||    || ||||||| ||||   || |
Db   1813 GTGATGTCAGTGGAAGATGAATCTACATC------CTGTTACTGCCTTTTGGACCCCTTT 1866

Qy   1867 GCCTGCTACGTCTTCACCGAGCAGCTGGGCCGCTTTGCCCTGGTGGGAGAGGCCCTCAGC 1926
          || ||  | ||  ||    || ||||| | ||  || ||| ||||||||| |   |||
Db   1867 GCGTGTCATGTGCTCCTGGACAGCTTTGGGACCTATGCGCTCACTGGAGAGCCAATCACA 1926

Qy   1927 GTGGCTGCCGCCAAGCGCCTCAAGCTGCTTCTGTTTGCGCCGGTGGCCTGCACCTCCCTC 1986
          |    ||||| ||||   || ||| || |  ||||     || |||| |||||||
Db   1927 GACTGTGCCGTGAAGCAACTGAAGGTGGCGGTTTTTGGCTGCATGTCCTGTAACTCCCTG 1986

Qy   1987 GAGTACAACATCCGGGTCTACTGCCTGCATGACACCCACGATGCACTCAAGGAGGTGGTG 2046
          || |||||| |  | || ||||| ||  |  | |||| ||||  |   |||| |||||
Db   1987 GATTACAACTTGAGAGTTTACTGTGTGGACAATACCCCTTGTGCATTTCAGGAAGTGGTT 2046

Qy   2047 CAGCTGGAGAAGCAGCTGGGGGGACAGCTGATCCAGGAGCCACGGGTCCTGCACTTCAAG 2106
           || | ||| | ||| |||||||| | | || ||| ` | ||||| |||||
Db   2047 TCAGATGAAAGGCATCAAGGTGGACAGCTCCTGGAAGAACCAAAATTGCTGCATTTCAAA 2106

Qy   2107 GACAGTTACCACAACCTGCGCCTATCCATCCACGATGTGCCCAGCTCCCTGTGGAAGAGT 2166
          |  |  | |    |   || |  | || |  |||  ||| | ||  | | ||||| | |
Db   2107 GGGAATACCTTTAGTCTTCAGATTTCTGTCCTTGATATTCCCCCATTCCTCTGGAGAATT 2166

Qy   2167 AAGCTCCTTGTCAGCTACCAGGAGATCCCCTTTTATCACATCTGGAATGGCACGCAGCGG 2226
          || |    |     || |||||| |||| || |   | || |   || | | |  || |
Db   2167 AAACCATTCACTGCCTGCCAGGAAGTCCCGTTCTCCCGCGTGTGGTGCAGTAACCGGCAG 2226

Qy   2227 TACTTGCACTGCACCTTCACCCTGGAGCGTGTCAGCCCCAGCACTAGTGACCTGGCCTGC 2286
           | ||||||| |||||  ||||| ||||||||||||||||| | |  |   | ||| |||||
Db   2227 CCCCTGCACTGTGCCTTCTCCCTGGAGCGTTATACGCCCACTACCACCCAGCTGTCCTGC 2286

Qy   2287 AAGCTGTGGGTGTGGCAGGTGGAGGGCGACGGGCAGAGCTTCAGCATCAACTTCAACATC 2346
          ||   | ||   |  ||||| |  | |||| | |  ||||  | ||   | |       |||
Db   2287 AAAATCTGCATTCGGCAGCTCAAAGGCCATGAACAGATCCTCCAAGTGCAGACATCAATC 2346

Qy   2347 ACCAAGGACACAAGGTTTGCTGAGCTGCTGGCTCTGGAGAGTGAAGCGGGGGTCCCAGCC 2406
           ||     | |  | |    |        |      |    |  ||     |||| ||
Db   2347 CTAGAGAGTGAACGAGAAACCATCACTTTCTTCGCACAAGAGGACAGCACTTTCCCTGCA 2406
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Qy     2407 CTGGTGGGCCCCAGTGCCTTCAAGATCCCCTTCCTCATTCGGCAGAAGATAATTTCCAGC 2466
             | |   ||||-|||  |||||||| || |||| |  ||| | ||| ||| | ||
Db     2407 CAGACTGGCCCCAAAGCCTTCAAAATTCCCTACTCCATCAGACAGCGGATTTGTGCTACA 2466


Qy     2467 CTGGACCCACCCTGTAGGCGGGGTGCCGACTGGCGGACTCTGGCCCAGAAACTCCACCTG 2526
             · | ||  | |||  |      ||    |||||||| ||  | || |||||| |  | |
Db     2467 TTTGATACCCCCAATGCCAAAGGCAAGGACTGGCAGATGTTAGCACAGAAAAACAGCATC 2526


Qy     2527 GACAGCCATCTCAGCTTCTTTGCCTCCAAGCCCAGCCCCACAGCCATGATCCTCAACCTG 2586
             ||||  || |      |  || ||  | |     ||||| | || | ||  | ||||||
Db     2527 AACAGGAATTTATCTTATTTCGCTACACAAAGTAGCCCATCTGCTGTCATTTTGAACCTG 2586


Qy     2587 TGGGAGGCGCGGCACTTCCCCAACGGCAACCTCAGCCAGCTGGCTGCAGCAGTGGCTGGA 2646
             |||||  || || ||     |  | ||  | ||    |     |||||   || | | |
Db     2587 TGGGAAGCTCGTCATCAGCATGATGGTGATCTTGACTCCCTGGCCTGTGCCCTTGAAGAG 2646


Qy     2647 CTGGGCCAGCCAGACGCTGGCCTCTTCACAGTGTCGGAGGCTGAG 2691
             | ||   | || || |    |||| |  | || || | ||
Db     2647 ATTGGGAGGACACACACGAAACTCTCAAACATTTCAGAATCCCAG 2691



RESULT 9
US-09-969-532-33
; Sequence 33, Application US/09969532
; Patent No. 6777232
; GENERAL INFORMATION:
;  APPLICANT: Walke, D. Wade
;  APPLICANT:  Scoville, John
;  TITLE OF INVENTION: No. 6777232el Human Membrane Proteins and Polynucleotides
Encoding the Same
;  FILE REFERENCE: LEX-0244-USA
;  CURRENT APPLICATION NUMBER: US/09/969,532
;  CURRENT FILING DATE:  2001-10-02
;  PRIOR APPLICATION NUMBER: US 60/237,280
;  PRIOR FILING DATE: 2000-10-02
;  NUMBER OF SEQ ID NOS: 33
;  SOFTWARE: FastSEQ for Windows Version 4.0
; SEQ ID NO 33
;   LENGTH: 3411
;   TYPE: DNA
;   ORGANISM: homo sapiens
US-09-969-532-33

  Query Match              18.1%;   Score 487;  DB 4;  Length 3411;
  Best Local Similarity    50.9%;   Pred. No. 4.4e-101;
  Matches 1305;  Conservative    0;  Mismatches 1230;  Indels   30;  Gaps    5;


Qy      127 CTTCCCCACTTCCTGGTGGAGCCCGAGGATGTGTACATCGTCAAGAACAAGCCAGTGCTG 186
             || || || ||| | | |||||| || |||| || || |||||| ||| || |
Db      271 CTGCCTCATTTCATAGAGGAGCCAGATGATGCTTATATTATCAAGAGCAACCCTATTGCA 330


Qy      187 CTTGTGTGCAAGGCCGTGCCCGCCACGCAGATCTTCTTCAAGTGCAACGGGGAGTGGGTG 246
             ||    ||||||| ||    ||| |||| ||||||| ||||||||| |||||||| |||||||||
Db      331 CTCAGGTGCAAAGCGAGGCCAGCCATGCAGATATTCTTCAAATGCAACGGCGAGTGGGTC 390


Qy      247 CGCCAGGTGGACCACGTGATCGAGCGCAGCACAGACGGGAGCAGTGGGCTGCCCACCATG 306
             |  ||| || |||||    ||  |    |||| ||||  || || |
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Db    391 CATCAGAACGAGCACGTCTCTGAAGAGACTCTGGACGAGAGCTCAGGTTTGAAGGTCCGC 450

Qy    307 GAGGTCCGCATTAATGTCTCAAGGCAGCAGGTCGAGAAGGTGTTCGGGCTGGAGGAATAC 366
              ||  ||    |||  |||||   |  | |||||  |||||  |||  |  |    ||||  |||||  ||
Db    451 GAAGTGTTCATCAATGTTACTAGGCAACAGGTGGAGGACTTCCATGGGCCCGAGGACTAT 510

Qy   .367 TGGTGCCAGTGCGTGGCATGGAGCTCCTCGGGCACCACCAAGAGTCAGAAGGCCTACATC 426
              ||||||||||| |||||  ||||||   |   |||  |||.|||||||   ||||||||    |
Db    511 TGGTGCCAGTGTGTGGCGTGGAGCCACCTGGGTACCTCCAAGAGCAGGAAGGCCTCTGTG 570

Qy    427 CGCATAGCCAGATTGCGCAAGAACTTCGAGCAGGAGCCGCTGGCCAAGGAGGTGTCCCTG 486
              |||||||||    || || ||  |||||  || || || || ||  |  |  |  ||| ||  || |
Db    571 CGCATAGCCTATTTACGGAAAAACTTTGAACAAGACCCACAAGGAAGGGAAGTTCCCATT 630

Qy    487 GAGCAGGGCATCGTGCTGCCCTGCCGTCCACCGGAGGGCATCCCTCCAGCCGAGGTGGAG 546
              ||          || ||  |||| ||||||  |||||  ||||| |||||.| ||||||||||||
Db    631 GAAGGCATGATTGTACTGCACTGCCGCCCACCAGAGGGAGTCCCTGCTGCCGAGGTGGAA 690

Qy    547 TGGCTCCGGAACGAGGACCTGGTGGACCCGTCCCTGGACCCCAATGTATACATCACGCGG 606
              |||||     || || || || |    | ||| |    |  |||    ||  |   ||| || |
Db    691 TGGCTGAAAAATGAAGAGCCCATTGACTCTGAACAAGACGAGAACATTGACACCAGGGCT 750

Qy    607 GAGCACAGCCTGGTGGTGCGACAGGCCCGCCTTGCTGACACGGCCAACTACACCTGCGTG 666
              || || | |||| |   | ||||||  || ||  | |||  |  |  || ||||||||||| ||
Db    751 GACCATAACCTGATCATCAGGCAGGCACGGCTCTCGGACTCAGGAAATTACACCTGCATG 810

Qy    667 GCCAAGAACATCGTGGCACGTCGCCGCAGCGCCTCCGCTGCTGTCATCGTCTACGTGAAC 726
              ||   |||||||||||    | | |||  || || || ||||  | ||||||||| |
Db    811 GCAGCCAACATCGTGGCTAAGAGGAGAAGCCTGTCGGCCACTGTTGTGGTCTACGTGGAT 870

Qy    727 GGTGGGTGGTCGACGTGGACCGAGTGGTCCGTCTGCAGCGCCAGCTGTGGGCGCGGCTGG 786
              ||  |  |||     ||||| |||  |||||||||||||||||    |      ||||
Db    871 GGGAGCTGGGAAGTGTGGAGCGAATGGTCCGTCTGCAGTCCAGAGTGTGA---------A 921

Qy    787 CAGAAACGGAGCCGGAGCTGCACCAACCCGGCGCCTCTCAACGGGGGCGCTTTCTGTGAG 846
              ||     |||| |||| |||||     ||  | |  ||||| ||||||  |||||||
Db    922 CATTTGCGGATCCGGGAGTGCACAGCACCACCCCCGAGAAATGGGGGCAAATTCTGTGAA 981

Qy    847 GGGCAGAATGTCCAGAAAACAGCCTGCGCCACCCTGTGCCCAGTAGACGGCAGCTGGAGC 906
              || | |   |        | | ||  |  |||| |       |   |  | |
Db    982 GGTCTAAGCCAGGAATCTGAAAACTGCACAGATGGTCTTTGCATCCTAGATAAAAAACCT 1041

Qy    907 CCGTGGAGCAAGTGGTCGGCCTGTGGGCTGGACTGCACCCACTGGCGGAGCCGTGAGTGC 966
              |        |       |       |  | ||  || | ||             | ||
Db   1042 CTTCATGAAATAAAACCCCAAAGCATTGAGAATGCCAGCGACATTGCTTTGTACTCGGGC 1101

Qy    967 TCTGACCCAGCACCCCGCAACGGAGGGGAGGAGTGCCAGGGCACTGACCTGGACACCCGC 1026
              | |   | ||  |      ||  | |   |   | |    ||   ||| |||  ||
Db   1102 TTGGGTGCTGCCGTCGTGGCCGTTGCAGTCCTGGTCATTGGTGTCACCCTTTACAGACGG 1161

Qy   1027 AACTGTACCAGTGACCTCTGTGTACACAGTGCTTCTGGCCCTGAGGACGTGGCCCTCTAT 1086
              | |   |     |   |   |   || || |||     |   |   | ||| |||
Db   1162 AGCCAGAGTGACTATGGCGTGGACGTCATTGACTCTTCTGCATTGACAGGTGGCTTCCA- 1220

Qy   1087 GTGGGCCTCATCGCCGTGGCCGTCTGCCTGGTCCTGCTGCTGCTTGTCCTCATCCTCGTT 1146
              | | |||  | |   | ||| |  | | |  |   |   ||  | ||  |  || || |
Db   1221 --GACCTTCAACTTCAAAACAGTCCGTCAAGCCAAGAATATCATGGAACTAATGATACAA 1278
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Qy   1147 TATTGCCGGAAGAAGGAGGGGCTGGACTCAGATGTGGCTGACTCGTCCATTCTCACCTCA 1206
               |           |   |||  |    ||  ||      | |    ||| ||
Db   1279 GAAAAATCCTTTGGTAACTCCCTGCTCCTGAATTCTGCCATGCAGCCAGATCTGACAGTG 1338

Qy   1207 GGCTTCCAGCCCGTCAGCATCAAGCCCAGCAAAGCAGACAACCCCCATCTGCTCACCATC 1266
               ||       |   | |    ||           | |       | |    ||||| |
Db   1339 AGCCGGACATACAGCGGACCCATCTGTCTGCAGGACCCTCTGGACAAGGAGCTCATGACA 1398

Qy   1267 CAGCCGGACCTCAGCACCACCACCACCACCTACCAGGGCAGTCTCTGTCCCCGGCAGGAT 1326
               || |   |||   | |  |    |   | | | | |  | ||    | |
Db   1399 GAGTCCTCACTCTTTAACCCTTTGTCGGACATCAAAGTGAAAGTCCAGAGCTCGTTCATG 1458

Qy   1327 GGGCCCAGCCCCAAGTTCCAGCTCACCAATGGGCACCTGCTCAGCCCCCTGGGTGGCGGC 1386
               |   ||       ||   ||    | |   |||       | |       |
Db   1459 GTTTCCCTGGGAGTGTCTGAGAGAGCTGAGTACCACGGCAAGAATCATTCCAGGACTTTT 1518

Qy   1387 CGCCACACACTGCACCACAGCTCTCCCACCTCTGAGGCCGAGGAGTTCGTCTCCCGCCTC 1446
               | |||    |     ||||||||| |   ||       |   ||      |   |  | | | ||
Db   1519 CCCCATGGAAACAACCACAGCTTTAGTACAATGCATCCCAGAAATAAAATGCCCTACATC 1578

Qy   1447 TCCACCCAGAACTACTTCCGCTCCCTGCCCCGAGGCACCAGCAACATGACCTATGGGACC 1506
               |   |  |              ||  ||  |||   | |  | |      |  |||    |||     |
Db   1579 CAAAATCTGTC---------ATCACTCCCCACAAGGACAGAACTGAGGACAACTGGTGTC 1629

Qy   1507 TTCAACTTCCTCGGGGGCCGGCTGATGATCCCTAATACAGGTATCAGCCTCCTCATCCCC 1566
               ||    |      |  |||||  ||  |   | ||  || |||||||| |  | |||  | ||||||  ||
Db   1630 TTTGGCCATTTAGGGGGGCGCTTAGTAATGCCAAATACAGGGGTGAGCTTACTCATACCA 1689

Qy   1567 CCAGATGCCATACCCCGAGGGAAGATCTATGAGATCTACCTCACGCTGCACAAGCCGGAA 1626
               |  | |||||||  ||      |  |||        |  |||||  ||  |   |   |     |||  |  |
Db   1690 CACGGTGCCATCCCAGAGGAGAATTCTTGGGAGATTTATATGTCCAT---CAACCAAGGT 1746

Qy   1627 GACGTGAGGTTGCCCCTAGCTGGCTGTCAGACCCTGCTGAGTCCCATCGTTAGCTGTGGA 1686
               ||       ||  | |        ||  |||||  |  | |     ||  |||||||||      || |  ||||||
Db   1747 GAACCCAGCCTCCAGTCAGATGGCTCTGAGGTGCTCCTGAGTCCTGAAGTCACCTGTGGT 1806

Qy   1687 CCCCCTGGCGTCCTGCTCACCCGGCCAGTCATCCTGGCTATGGACCACTGTGGGGAGCCC 1746
               || || | | |  |   |||||    ||  |      || | ||       |||||||    ||      |
Db   1807 CCTCCAGACATGATCGTCACCACTCCCTTTGCATTGACCATCCCGCACTGTGCAGATGTC 1866

Qy   1747 AGCCCTGACAGCTGGAGCCTGCGCCTCAAAAAGCAGTCGTGCGAGGGCAGCTGGGAGGAT 1806
               ||  ||||       ||||    | |    | ||  |||    | |       ||||||    ||||||||
Db   1867 AGTTCTGAGCATTGGAATATCCATTTAAAGAAGAGGACACAGCAGGGCAAATGGGAGGAA 1926

Qy   1807 GTGCTGCACCTGGGCGAGGAGGCGCCCTCCCACCTCTACTACTGCCAGCTGGAGGCCAGT 1866
               ||| ||    |||   || ||    |   | |          ||   ||||||||     ||||    || |
Db   1927 GTGATGTCAGTGGAAGATGAATCTACATC------CTGTTACTGCCTTTTGGACCCCTTT 1980

Qy   1867 GCCTGCTACGTCTTCACCGAGCAGCTGGGCCGCTTTGCCCTGGTGGGAGAGGCCCTCAGC 1926
               || ||  | ||  ||   ||     | ||    | ||  || |||  ||    ||||||| |    |||
Db   1981 GCGTGTCATGTGCTCCTGGACAGCTTTGGGACCTATGCGCTCACTGGAGAGCCAATCACA 2040

Qy   1927 GTGGCTGCCGCCAAGCGCCTCAAGCTGCTTCTGTTTGCGCCGGTGGCCTGCACCTCCCTC 1986
               |     |||||  ||||   || ||| ||      | ||||          || ||||| |  ||||||
Db   2041 GACTGTGCCGTGAAGCAACTGAAGGTGGCGGTTTTTGGCTGCATGTCCTGTAACTCCCTG 2100
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Qy    1987 GAGTACAACATCCGGGTCTACTGCCTGCATGACACCCACGATGCACTCAAGGAGGTGGTG 2046
           ||  |||||| |   | || ||||| || |   | ||||    |||| |  |||| |||||
Db    2101 GATTACAACTTGAGAGTTTACTGTGTGGACAATACCCCTTGTGCATTTCAGGAAGTGGTT 2160

Qy    2047 CAGCTGGAGAAGCAGCTGGGGGGACAGCTGATCCAGGAGCCACGGGTCCTGCACTTCAAG 2106
           || | ||| |  || ||||||||| |   | || |||    | |||||| |||||
Db    2161 TCAGATGAAAGGCATCAAGGTGGACAGCTCCTGGAAGAACCAAAATTGCTGCATTTCAAA 2220

Qy    2107 GACAGTTACCACAACCTGCGCCTATCCATCCACGATGTGCCCAGCTCCCTGTGGAAGAGT 2166
           |  | |  |   |  ||  |   | || |||  ||| | |||  | ||| ||||  | |
Db    2221 GGGAATACCTTTAGTCTTCAGATTTCTGTCCTTGATATTCCCCCATTCCTCTGGAGAATT 2280

Qy    2167 AAGCTCCTTGTCAGCTACCAGGAGATCCCCTTTTATCACATCTGGAATGGCACGCAGCGG 2226
           || |   |      || |||||| |||| || |   | |  | |||    | |  | ||  |
Db    2281 AAACCATTCACTGCCTGCCAGGAAGTCCCGTTCTCCCGCGTGTGGTGCAGTAACCGGCAG 2340

Qy    2227 TACTTGCACTGCACCTTCACCCTGGAGCGTGTCAGCCCCAGCACTAGTGACCTGGCCTGC 2286
           | |||||||| ||||| |||||||||||| |   | |||| || |   | ||| |||||
Db    2341 CCCCTGCACTGTGCCTTCTCCCTGGAGCGTTATACGCCCACTACCACCCAGCTGTCCTGC 2400

Qy    2287 AAGCTGTGGGTGTGGCAGGTGGAGGGCGACGGGCAGAGCTTCAGCATCAACTTCAACATC 2346
           ||   | ||  | |||||| |  | ||| | |  |||| | ||         |   |||
Db    2401 AAAATCTGCATTCGGCAGCTCAAAGGCCATGAACAGATCCTCCAAGTGCAGACATCAATC 2460

Qy    2347 ACCAAGGACACAAGGTTTGCTGAGCTGCTGGCTCTGGAGAGTGAAGCGGGGGTCCCAGCC 2406
           ||      | |     |         |        |    |   ||    |||| ||
Db    2461 CTAGAGAGTGAACGAGAAACCATCACTTTCTTCGCACAAGAGGACAGCACTTTCCCTGCA 2520

Qy    2407 CTGGTGGGCCCCAGTGCCTTCAAGATCCCCTTCCTCATTCGGCAGAAGATAATTTCCAGC 2466
           | |   |||||||  |||||||| ||  ||||  |   ||| | ||| |||   |  | |
Db    2521 CAGACTGGCCCCAAAGCCTTCAAAATTCCCTACTCCATCAGACAGCGGATTTGTGCTACA 2580

Qy    2467 CTGGACCCACCCTGTAGGCGGGGTGCCGACTGGCGGACTCTGGCCCAGAAACTCCACCTG 2526
           | || | |||  |         ||       |||||| ||  | || ||||||  |  | |
Db    2581 TTTGATACCCCCAATGCCAAAGGCAAGGACTGGCAGATGTTAGCACAGAAAAACAGCATC 2640

Qy    2527 GACAGCCATCTCAGCTTCTTTGCCTCCAAGCCCAGCCCCACAGCCATGATCCTCAACCTG 2586
           ||||  || |    |  | || ||  |    |||||  | ||  | || | |||||||
Db    2641 AACAGGAATTTATCTTATTTCGCTACACAAAGTAGCCCATCTGCTGTCATTTTGAACCTG 2700

Qy    2587 TGGGAGGCGCGGCACTTCCCCAACGGCAACCTCAGCCAGCTGGCTGCAGCAGTGGCTGGA 2646
           ||||| || || || ||     |   | || | ||   |     ||||||   || | | |
Db    2701 TGGGAAGCTCGTCATCAGCATGATGGTGATCTTGACTCCCTGGCCTGTGCCCTTGAAGAG 2760

Qy    2647 CTGGGCCAGCCAGACGCTGGCCTCTTCACAGTGTCGGAGGCTGAG 2691
           | ||   | || || | |    |||| |   | || || | ||
Db    2761 ATTGGGAGGACACACACGAAACTCTCAAACATTTCAGAATCCCAG 2805
```

RESULT 10
US-09-969-532-11
; Sequence 11, Application US/09969532
; Patent No. 6777232
; GENERAL INFORMATION:
; APPLICANT: Walke, D. Wade
; APPLICANT: Scoville, John

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  Query Match              17.3%;   Score 467;  DB 4;  Length 2703;
  Best Local Similarity    51.0%;   Pred. No. 1.5e-96;
  Matches 1307;  Conservative    0;  Mismatches 1195;  Indels  63;  Gaps    6;


Qy       127 CTTCCCCACTTCCTGGTGGAGCCCGAGGATGTGTACATCGTCAAGAACAAGCCAGTGCTG 186
             || || || ||| | | |||||| || |||| || || |||||| ||| || |
Db       157 CTGCCTCATTTCATAGAGGAGCCAGATGATGCTTATATTATCAAGAGCAACCCTATTGCA 216


Qy       187 CTTGTGTGCAAGGCCGTGCCCGCCACGCAGATCTTCTTCAAGTGCAACGGGGAGTGGGTG 246
             ||    |||||| ||    ||| |||| |||||| ||||||||| ||||||||| |||||||||
Db       217 CTCAGGTGCAAAGCGAGGCCAGCCATGCAGATATTCTTCAAATGCAACGGCGAGTGGGTC 276


Qy       247 CGCCAGGTGGACCACGTGATCGAGCGCAGCACAGACGGGAGCAGTGGGCTGCCCACCATG 306
             | ||| || ||||| || | |||| |||| || || |
Db       277 CATCAGAACGAGCACGTCTCTGAAGAGACTCTGGACGAGAGCTCAGGTTTGAAGGTCCGC 336


Qy       307 GAGGTCCGCATTAATGTCTCAAGGCAGCAGGTCGAGAAGGTGTTCGGGCTGGAGGAATAC 366
             || || ||| ||||| | ||||| ||||| ||| | | |||| ||||| ||
Db       337 GAAGTGTTCATCAATGTTACTAGGCAACAGGTGGAGGACTTCCATGGGCCCGAGGACTAT 396


Qy       367 TGGTGCCAGTGCGTGGCATGGAGCTCCTCGGGCACCACCAAGAGTCAGAAGGCCTACATC 426
             |||||||||||| ||||| |||||| | ||| ||| ||||||| |||||||| |
Db       397 TGGTGCCAGTGTGTGGCGTGGAGCCACCTGGGTACCTCCAAGAGCAGGAAGGCCTCTGTG 456


Qy       427 CGCATAGCCAGATTGCGCAAGAACTTCGAGCAGGAGCCGCTGGCCAAGGAGGTGTCCCTG 486
             |||||||||   || || || ||||| || || || || ||-|  |  | ||| ||  || |
Db       457 CGCATAGCCTATTTACGGAAAAACTTTGAACAAGACCCACAAGGAAGGGAAGTTCCCATT 516


Qy       487 GAGCAGGGCATCGTGCTGCCCTGCCGTCCACCGGAGGGCATCCCTCCAGCCGAGGTGGAG 546
             ||        || || |||| ||||||| ||||| ||||| |||||| | |||||||||||||
Db       517 GAAGGCATGATTGTACTGCACTGCCGCCCACCAGAGGGAGTCCCTGCTGCCGAGGTGGAA 576


Qy       547 TGGCTCCGGAACGAGGACCTGGTGGACCCGTCCCTGGACCCCAATGTATACATCACGCGG 606
             |||||     || || || ||  | | ||| |   | |||   || | ||| || |
Db       577 TGGCTGAAAAATGAAGAGCCCATTGACTCTGAACAAGACGAGAACATTGACACCAGGGCT 636


Qy       607 GAGCACAGCCTGGTGGTGCGACAGGCCCGCCTTGCTGACACGGCCAACTACACCTGCGTG 666
             || || | |||| | | | |||||| || || | ||| | | || |||||||||| ||
Db       637 GACCATAACCTGATCATCAGGCAGGCACGGCTCTCGGACTCAGGAAATTACACCTGCATG 696


Qy       667 GCCAAGAACATCGTGGCACGTCGCCGCAGCGCCTCCGCTGCTGTCATCGTCTACGTGAAC 726
             ||       ||||||||||||        |  | |||  || || |||| | ||||||||| |
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Db    697 GCAGCCAACATCGTGGCTAAGAGGAGAAGCCTGTCGGCCACTGTTGTGGTCTACGTGGAT 756

Qy    727 GGTGGGTGGTCGACGTGGACCGAGTGGTCCGTCTGCAGCGCCAGCTGTGGGCGCGGCTGG 786
              ||   | |||     ||||| |||  ||||||||||||||||  |     ||||
Db    757 GGGAGCTGGGAAGTGTGGAGCGAATGGTCCGTCTGCAGTCCAGAGTGTG---------AA 807

Qy    787 CAGAAACGGAGCCGGAGCTGCACCAACCCGGCGCCTCTCAACGGGGGCGCTTTCTGTGAG 846
              ||    |||| ||||     |||||    || | ||    || ||||||   ||||||||
Db    808 CATTTGCGGATCCGGGAGTGCACAGCACCACCCCCGAGAAATGGGGGCAAATTCTGTGAA 867

Qy    847 GGGCAGAATGTCCAGAAAACAGCCTGCGCCACCCTGTGCCCAGTAGACGGCAGCTGGAGC 906
              || | |        |      |  |||| |        |            |||| ||
Db    868 GGTCTAAGCCAGGAATCTGAAAACTGCACAGATGGTCTTTGCATCCTAGGCATTGAGAAT 927

Qy    907 CCGTGGAGCAAGTGGTCGGCCTGTGGGCTGGACTGCACCCACTGGCGGAGCCGTGAGTGC 966
               | |  ||    | |   || ||  ||  |||                         ||||
Db    928 GCCAGCGACATTGCTTTGTACTCGGGCTTGG----------------------GTGC 962

Qy    967 TCTGACCCAGCACCCCGCAACGGAGGGGAGGAGTGCCAGGGCACTGACCTGGACACCCGC 1026
              |     |  |  |   |||      ||  |   ||| ||       |      |   |||
Db    963 TGCCGTCGTGGCCGTTGCAGTCCTGGTCATTGGTGTCACCCTTTACAGACGGA------- 1015

Qy   1027 AACTGTACCAGTGACCTCTGTGTACACAGTGCTTCTGGCCCTGAGGACGTGGCCCTCTAT 1086
               |     |||||| |  ||   ||       |   |||           |   | ||
Db   1016 ----GCCAGAGTGACTATGGCGTGGACGTCATTGACTCTTCTGCATTGACAGGTGGCTTC 1071

Qy   1087 GTGGGCCTCATCGCCGTGGCCGTCTGCCTGGTCCTGCTGCTGCTTGTCCTCATCCTCGTT 1146
               | |  ||| |  |  | ||| |  | |   |  |  |     | | |  || || |
Db   1072 CAGACCTTCAACTTCAAAACAGTCCGTCAAGCCAAGAATATCATGGAACTAATGATACAA 1131

Qy   1147 TATTGCCGGAAGAAGGAGGGGCTGGACTCAGATGTGGCTGACTCGTCCATTCTCACCTCA 1206
               |        |      |   |||  |   |   ||   ||      | |   ||| ||
Db   1132 GAAAAATCCTTTGGTAACTCCCTGCTCCTGAATTCTGCCATGCAGCCAGATCTGACAGTG 1191

Qy   1207 GGCTTCCAGCCCGTCAGCATCAAGCCCAGCAAAGCAGACAACCCCCATCTGCTCACCATC 1266
               ||         | | |   ||         | |          | |   |||||  |
Db   1192 AGCCGGACATACAGCGGACCCATCTGTCTGCAGGACCCTCTGGACAAGGAGCTCATGACA 1251

Qy   1267 CAGCCGGACCTCAGCACCACCACCACCACCTACCAGGGCAGTCTCTGTCCCCGGCAGGAT 1326
               || |     |||   |||      |   | |   |  ||   |      |
Db   1252 GAGTCCTCACTCTTTAACCCTTTGTCGGACATCAAAGTGAAAGTCCAGAGCTCGTTCATG 1311

Qy   1327 GGGCCCAGCCCCAAGTTCCAGCTCACCAATGGGCACCTGCTCAGCCCCCTGGGTGGCGGC 1386
               |   ||       ||   ||   | |   |||       | |       |
Db   1312 GTTTCCCTGGGAGTGTCTGAGAGAGCTGAGTACCACGGCAAGAATCATTCCAGGACTTTT 1371

Qy   1387 CGCCACACACTGCACCACAGCTCTCCCACCTCTGAGGCCGAGGAGTTCGTCTCCCGCCTC 1446
               | |||   |   ||||||||| |    ||      |   | |     |   ||  | ||
Db   1372 CCCCATGGAAACAACCACAGCTTTAGTACAATGCATCCCAGAAATAAAATGCCCTACATC 1431

Qy   1447 TCCACCCAGAACTACTTCCGCTCCCTGCCCCGAGGCACCAGCAACATGACCTATGGGACC 1506
               |   |       || || |||    | | ||    | |||    ||| |   |||  |
Db   1432 CAAAATCTGTC---------ATCACTCCCCACAAGGACAGAACTGAGGACAACTGGTGTC 1482

Qy   1507 TTCAACTTCCTCGGGGGCCGGCTGATGATCCCTAATACAGGTATCAGCCTCCTCATCCCC 1566
              ||   |     | ||||| ||  |  | | || || |||||||| | ||| |   ||||| ||
Db   1483 TTTGGCCATTTAGGGGGGCGCTTAGTAATGCCAAATACAGGGGTGAGCTTACTCATACCA 1542
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Qy   1567 CCAGATGCCATACCCCGAGGGAAGATCTATGAGATCTACCTCACGCTGCACAAGCCGGAA 1626
               |  | |||||| ||    | |||    |  | ||||| ||  |  |  |    ||| | |
Db   1543 CACGGTGCCATCCCAGAGGAGAATTCTTGGGAGATTTATATGTCCAT---CAACCAAGGT 1599

Qy   1627 GACGTGAGGTTGCCCCTAGCTGGCTGTCAGACCCTGCTGAGTCCCATCGTTAGCTGTGGA 1686
               ||     ||  | |      ||  |||||| | ||     || |||||||||    || | ||||||
Db   1600 GAACCCAGCCTCCAGTCAGATGGCTCTGAGGTGCTCCTGAGTCCTGAAGTCACCTGTGGT 1659

Qy   1687 CCCCCTGGCGTCCTGCTCACCCGGCCAGTCATCCTGGCTATGGACCACTGTGGGGAGCCC 1746
               || ||  |  | |    | |||||      ||  |       || | ||     |||||||  ||    |
Db   1660 CCTCCAGACATGATCGTCACCACTCCCTTTGCATTGACCATCCCGCACTGTGCAGATGTC 1719

Qy   1747 AGCCCTGACAGCTGGAGCCTGCGCCTCAAAAAGCAGTCGTGCGAGGGCAGCTGGGAGGAT 1806
               ||  ||||      |||| |    | |    | || |||   | |      ||||||  |||||||||
Db   1720 AGTTCTGAGCATTGGAATATCCATTTAAAGAAGAGGACACAGCAGGGCAAATGGGAGGAA 1779

Qy   1807 GTGCTGCACCTGGGCGAGGAGGCGCCCTCCCACCTCTACTACTGCCAGCTGGAGGCCAGT 1866
               ||| ||   ||| || || |  ||| ||     || ||||||| |||| || |
Db   1780 GTGATGTCAGTGGAAGATGAATCTACATC------CTGTTACTGCCTTTTGGACCCCTTT 1833

Qy   1867 GCCTGCTACGTCTTCACCGAGCAGCTGGGCCGCTTTGCCCTGGTGGGAGAGGCCCTCAGC 1926
               || ||  | ||  ||    ||     | ||  || || ||| ||    |||||| |   |||
Db   1834 GCGTGTCATGTGCTCCTGGACAGCTTTGGGACCTATGCGCTCACTGGAGAGCCAATCACA 1893

Qy   1927 GTGGCTGCCGCCAAGCGCCTCAAGCTGCTTCTGTTTGCGCCGGTGGCCTGCACCTCCCTC 1986
               |     ||||| ||||    || |||  ||   ||    | ||||      || |||| | ||||||
Db   1894 GACTGTGCCGTGAAGCAACTGAAGGTGGCGGTTTTTGGCTGCATGTCCTGTAACTCCCTG 1953

Qy   1987 GAGTACAACATCCGGGTCTACTGCCTGCATGACACCCACGATGCACTCAAGGAGGTGGTG 2046
               || |||||| |   | || ||||| ||  |  | ||||     |||| |   |||| |||||
Db   1954 GATTACAACTTGAGAGTTTACTGTGTGGACAATACCCCTTGTGCATTTCAGGAAGTGGTT 2013

Qy   2047 CAGCTGGAGAAGCAGCTGGGGGGACAGCTGATCCAGGAGCCACGGGTCCTGCACTTCAAG 2106
               || | ||| |    || ||||||||   |    | || |||     | ||||| |||||
Db   2014 TCAGATGAAAGGCATCAAGGTGGACAGCTCCTGGAAGAACCAAAATTGCTGCATTTCAAA 2073

Qy   2107 GACAGTTACCACAACCTGCGCCTATCCATCCACGATGTGCCCAGCTCCCTGTGGAAGAGT 2166
               |   | |  |  |    | ||  | | ||  |||  | | ||| |||  | ||| ||||  | |
Db   2074 GGGAATACCTTTAGTCTTCAGATTTCTGTCCTTGATATTCCCCCATTCCTCTGGAGAATT 2133

Qy   2167 AAGCTCCTTGTCAGCTACCAGGAGATCCCCTTTTATCACATCTGGAATGGCACGCAGCGG 2226
               || .|   |      || |||||| |||| || |  | | |  | |||   | |  | ||| |
Db   2134 AAACCATTCACTGCCTGCCAGGAAGTCCCGTTCTCCCGCGTGTGGTGCAGTAACCGGCAG 2193

Qy   2227 TACTTGCACTGCACCTTCACCCTGGAGCGTGTCAGCCCCAGCACTAGTGACCTGGCCTGC 2286
               |  |||||||| ||||| |||||||||||||||||||    | |||| || |   | ||| |||||
Db   2194 CCCCTGCACTGTGCCTTCTCCCTGGAGCGTTATACGCCCACTACCACCCAGCTGTCCTGC 2253

Qy   2287 AAGCTGTGGGTGTGGCAGGTGGAGGGCGACGGGCAGAGCTTCAGCATCAACTTCAACATC 2346
               ||  | | ||  |  ||||| |  | ||| | |  ||||  | ||    |  |         |||
Db   2254 AAAATCTGCATTCGGCAGCTCAAAGGCCATGAACAGATCCTCCAAGTGCAGACATCAATC 2313

Qy   2347 ACCAAGGACACAAGGTTTGCTGAGCTGCTGGCTCTGGAGAGTGAAGCGGGGGTCCCAGCC 2406
               ||    |  | |  |      |  |     |  |      |        ||      |||| ||
Db   2314 CTAGAGAGTGAACGAGAAACCATCACTTTCTTCGCACAAGAGGACAGCACTTTCCCTGCA 2373
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Qy    2407 CTGGTGGGCCCCAGTGCCTTCAAGATCCCCTTCCTCATTCGGCAGAAGATAATTTCCAGC 2466
                 | .|     |||||||.  |||||||| || ||||| |  ||| | ||| |||    | | |
Db    2374 CAGACTGGCCCCAAAGCCTTCAAAATTCCCTACTCCATCAGACAGCGGATTTGTGCTACA 2433

Qy    2467 CTGGACCCACCCTGTAGGCGGGGTGCCGACTGGCGGACTCTGGCCCAGAAACTCCACCTG 2526
                 | ||  | ||| |         ||      |||||||| ||   | || |||||| |  | |
Db    2434 TTTGATACCCCCAATGCCAAAGGCAAGGACTGGCAGATGTTAGCACAGAAAAACAGCATC 2493

Qy    2527 GACAGCCATCTCAGCTTCTTTGCCTCCAAGCCCAGCCCCACAGCCATGATCCTCAACCTG 2586
                 ||||  || |    |  || || | |      ||||| | || | || | ||||||
Db    2494 AACAGGAATTTATCTTATTTCGCTACACAAAGTAGCCCATCTGCTGTCATTTTGAACCTG 2553

Qy    2587 TGGGAGGCGCGGCACTTCCCCAACGGCAACCTCAGCCAGCTGGCTGCAGCAGTGGCTGGA 2646
                 ||||| || || ||      |   | || | ||   |    ||||||     || | | |
Db    2554 TGGGAAGCTCGTCATCAGCATGATGGTGATCTTGACTCCCTGGCCTGTGCCCTTGAAGAG 2613

Qy    2647 CTGGGCCAGCCAGACGCTGGCCTCTTCACAGTGTCGGAGGCTGAG 2691
                 |  ||   | || || ||  |    |||| |   | ||  || |  ||
Db    2614 ATTGGGAGGACACACACGAAACTCTCAAACATTTCAGAATCCCAG 2658


RESULT 11
US-09-969-532-13
; Sequence 13, Application US/09969532
; Patent No. 6777232
; GENERAL INFORMATION:
;  APPLICANT: Walke, D. Wade
;  APPLICANT:  Scoville, John
;  TITLE OF INVENTION: No. 6777232el Human Membrane Proteins and Polynucleotides
Encoding the Same
;  FILE REFERENCE: LEX-0244-USA
;  CURRENT APPLICATION NUMBER: US/09/969,532
;  CURRENT FILING DATE:  2001-10-02
;  PRIOR APPLICATION NUMBER: US 60/237,280
;  PRIOR FILING DATE: 2000-10-02
;  NUMBER OF SEQ ID NOS: 33
;  SOFTWARE: FastSEQ for Windows Version 4.0
; SEQ ID NO 13
;   LENGTH: 2694
;   TYPE: DNA
;   ORGANISM: homo sapiens
US-09-969-532-13

  Query Match            17.1%;  Score 460.2;  DB 4;  Length 2694;
  Best Local Similarity  51.9%;  Pred. No. 5.1e-95;
  Matches 1330;  Conservative   0;  Mismatches 1163;  Indels  72;  Gaps   10;

Qy     127 CTTCCCCACTTCCTGGTGGAGCCCGAGGATGTGTACATCGTCAAGAACAAGCCAGTGCTG 186
                 || || || ||| | | ||||||| || |||| || || |||||| ||| ||  |
Db     157 CTGCCTCATTTCATAGAGGAGCCAGATGATGCTTATATTATCAAGAGCAACCCTATTGCA 216

Qy     187 CTTGTGTGCAAGGCCGTGCCCGCCACGCAGATCTTCTTCAAGTGCAACGGGGAGTGGGTG 246
                 ||   |||||| ||   ||| |||| ||||||| ||||||||| |||||||| |||||||||
Db     217 CTCAGGTGCAAAGCGAGGCCAGCCATGCAGATATTCTTCAAATGCAACGGCGAGTGGGTC 276

Qy     247 CGCCAGGTGGACCACGTGATCGAGCGCAGCACAGACGGGAGCAGTGGGCTGCCCACCATG 306
                 |  ||| || ||||||    || |     |||| |||| || ||       |
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Db    277 CATCAGAACGAGCACGTCTCTGAAGAGACTCTGGACGAGAGCTCAGGTTTGAAGGTCCGC 336

Qy    307 GAGGTCCGCATTAATGTCTCAAGGCAGCAGGTCGAGAAGGTGTTCGGGCTGGAGGAATAC 366
              || ||   ||||  ||||||  |  |||||| |||||| |||  | |    |||| ||||| ||
Db    337 GAAGTGTTCATCAATGTTACTAGGCAACAGGTGGAGGACTTCCATGGGCCCGAGGACTAT 396

Qy    367 TGGTGCCAGTGCGTGGCATGGAGCTCCTCGGGCACCACCAAGAGTCAGAAGGCCTACATC 426
              |||||||||||| |||||| ||||||      |  ||| |||  |||||||    |||||||||    |
Db    397 TGGTGCCAGTGTGTGGCGTGGAGCCACCTGGGTACCTCCAAGAGCAGGAAGGCCTCTGTG 456

Qy    427 CGCATAGCCAGATTGCGCAAGAACTTCGAGCAGGAGCCGCTGGCCAAGGAGGTGTCCCTG 486
              ||||||||   ||  || || ||||| || || || || ||   |    | ||| ||  || |
Db    457 CGCATAGCCTATTTACGGAAAAACTTTGAACAAGACCCACAAGGAAGGGAAGTTCCCATT 516

Qy    487 GAGCAGGGCATCGTGCTGCCCTGCCGTCCACCGGAGGGCATCCCTCCAGCCGAGGTGGAG 546
              ||        || || |||| |||||| |||||| ||||||      |||||| | |||||||||||
Db    517 GAAGGCATGATTGTACTGCACTGCCGCCCACCAGAGGGAGTCCCTGCTGCCGAGGTGGAA 576

Qy    547 TGGCTCCGGAACGAGGACCTGGTGGACCCGTCCCTGGACCCCAATGTATACATCACGCGG 606
              |||||    || || || || |    | ||| |     |   ||| || | ||| || |
Db    577 TGGCTGAAAAATGAAGAGCCCATTGACTCTGAACAAGACGAGAACATTGACACCAGGGCT 636

Qy    607 GAGCACAGCCTGGTGGTGCGACAGGCCCGCCTTGCTGACACGGCCAACTACACCTGCGTG 666
              || || |  |||| |   | |  |||| || || |  || ||| | |  || |||||||||| ||
Db    637 GACCATAACCTGATCATCAGGCAGGCACGGCTCTCGGACTCAGGAAATTACACCTGCATG 696

Qy    667 GCCAAGAACATCGTGGCACGTCGCCGCAGCGCCTCCGCTGCTGTCATCGTCTACGTGAAC 726
              ||     |||||||||||     |   | |||    || || || |||| | ||||||||||| |
Db    697 GCAGCCAACATCGTGGCTAAGAGGAGAAGCCTGTCGGCCACTGTTGTGGTCTACGTGGAT 756

Qy    727 GGTGGGTGGTCGACGTGGACCGAGTGGTCCGTCTGCAGCGCCAGCTGTGGGCGCGGCTGG 786
              ||  |  |||     ||||| |||  ||||||||||||||||    |      ||||
Db    757 GGGAGCTGGGAAGTGTGGAGCGAATGGTCCGTCTGCAGTCCAGAGTGTG---------AA 807

Qy    787 CAGAAACGGAGCCGGAGCTGCACCAACCCGGCGCCTCTCAACGGGGGCGCTTTCTGTGAG 846
              ||     |||| ||||     ||||| |      || | ||    || ||||||    |||||||||
Db    808 CATTTGCGGATCCGGGAGTGCACAGCACCACCCCCGAGAAATGGGGGCAAATTCTGTGAA 867

Qy    847 GGGCAGAATGTCCAGAAAACAGCCTGCGCCACCCTGTGCCCAGTAGACGGCAGCTGGAGC 906
              || |   | | |   |||          ||      ||   | | | | |   |   | |
Db    868 GGTC--TAAGCCAGGAATCTGAAAACTGCACAGATGGTCTTTGCATCCTAGATAAAAAAC 925

Qy    907 CCGTGGAGCAAGTGGTCGGCCTGTGGGCTGGACTGCACCCACTGGCGGAGCCGTGAGTGC 966
              | | |  | ||     ||  | || |  || |   | |   |   | | | | | ||||
Db    926 CTCTTCATGAAATAAAACCCCAAAGCATTGAG----AATGCCAGCGACATTGCTTTGTAC 981

Qy    967 TCTGACCCAGCACCCCGCAACGGAGGGGAGGAGTGCCAGGGCACTGACCTGGACACCCGC 1026
              || | |   |   |  | |   |  |  |        || || || || ||      | |
Db    982 TCGGGCTTGGGTGCTGCCGTCGTGGCCGTTGCAGTCCTGGTCATTGGTGT------CACC 1035

Qy   1027 AACTGTACCAGTGACCTCTGTGTACACAGTGCTTCTGGCCCTGAGGACGTGGCCCTCTAT 1086
               |  |   |    || ||| | || | | |     | | | | ||      || |
Db   1036 CTTTACAGACGGAGCCAGAGTGACTATGGCGTGGACGTCATTGACTCTTCTGCATTGACA 1095

Qy   1087 GTGGGCCTCATCGCCGTGGCCGTCTGCCTGGTCCTGCTGCTGCTTGTCCTCATCCTCGTT 1146
               |  ||| ||   || |  || |  ||||    |     |  | |   ||| |||| |
Db   1096 GGTGGCTTCCAGACCTTCAACTTCAAAACAGTCCGTCAAGGTAACTCCCTGCTCCTGAAT 1155
```

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Qy   1147 TATTGCCGGAAGAAGGAGGGGCTGGACTCAGATGTGGCTGACTCGTCCATTCTCACCTCA 1206
          | | | | |        | | |  || | |   || |      |     | | ||
Db   1156 TCTGCCATGCA--------GCCAGATCTGACAGTGAGCCGGACATACAGCGGACCCATCT 1207

Qy   1207 GGCTTCCAGCCCGTCAGCATCAAGCCCAGCAAAGCAGACAACCCCCATCTGCTCACCATC 1266
          | || | | || || | | | |    ||| | | | | || ||    | ||| |
Db   1208 GTCTGCAGGACCCTCTGGACAAGG---AGCTCATGACAGAGTCCTCACTCTTTAACCCTT 1264

Qy   1267 CAGCCGGACCTCAGCACCACCACCACCACCTACCAGGGCAGTCTCTGTCCCCGGCAGGAT 1326
           | |    |      |    ||| |      |   || |||    | |
Db   1265 TGTCGGACATCAAAGTGAAAGTCCAGAGCTCGTTCATGGTTTCCCTGGGAGTGTCTGAGA 1324

Qy   1327 GGGCCCAGCCCCAAGTTCCAGCTCACCAATGGGCACCTGCTCAGCCCCCTGGGTGGCGGC 1386
          | || || ||| |       |||      || |    |
Db   1325 GAGCTGAGTACCACGGCAAGAATCATTCCAGGACTTTT---------------------- 1362

Qy   1387 CGCCACACACTGCACCACAGCTCTCCCACCTCTGAGGCCGAGGAGTTCGTCTCCCGCCTC 1446
          | ||| |     ||||||||| |   ||     | ||     |     | ||  | ||
Db   1363 CCCCATGGAAACAACCACAGCTTTAGTACAATGCATCCCAGAAATAAAATGCCCTACATC 1422

Qy   1447 TCCACCCAGAACTACTTCCGCTCCCTGCCCCGAGGCACCAGCAACATGACCTATGGGACC 1506
           |  | |          || || ||| | | ||     | ||| |||    |
Db   1423 CAAAATCTG---------TCATCACTCCCCACAAGGACAGAACTGAGGACAACTGGTGTC 1473

Qy   1507 TTCAACTTCCTCGGGGGCCGGCTGATGATCCCTAATACAGGTATCAGCCTCCTCATCCCC 1566
          ||   |     | |||||| ||  |   | ||  || ||||||||| | ||| | ||||| ||
Db   1474 TTTGGCCATTTAGGGGGGCGCTTAGTAATGCCAAATACAGGGGTGAGCTTACTCATACCA 1533

Qy   1567 CCAGATGCCATACCCCGAGGGAAGATCTATGAGATCTACCTCACGCTGCACAAGCCGGAA 1626
          | | |||||||| ||    | |||      | |||||| ||  | | |   ||| | |
Db   1534 CACGGTGCCATCCCAGAGGAGAATTCTTGGGAGATTTATATGTCCAT---CAACCAAGGT 1590

Qy   1627 GACGTGAGGTTGCCCCTAGCTGGCTGTCAGACCCTGCTGAGTCCCATCGTTAGCTGTGGA 1686
          ||     || | |    || ||||| | ||   || |||||||||    || | ||||||
Db   1591 GAACCCAGCCTCCAGTCAGATGGCTCTGAGGTGCTCCTGAGTCCTGAAGTCACCTGTGGT 1650

Qy   1687 CCCCCTGGCGTCCTGCTCACCCGGCCAGTCATCCTGGCTATGGACCACTGTGGGGAGCCC 1746
           || || | | |   | | ||| | ||||| ||   | | | || |||||||| ||   |
Db   1651 CCTCCAGACATGATCGTCACCACTCCCTTTGCATTGACCATCCCGCACTGTGCAGATGTC 1710

Qy   1747 AGCCCTGACAGCTGGAGCCTGCGCCTCAAAAAGCAGTCGTGCGAGGGCAGCTGGGAGGAT 1806
          ||  |||    ||||   | |  |  | || ||| | |   |||||| ||||||||||||||
Db   1711 AGTTCTGAGCATTGGAATATCCATTTAAAGAAGAGGACACAGCAGGGCAAATGGGAGGAA 1770

Qy   1807 GTGCTGCACCTGGGCGAGGAGGCGCCCTCCCACCTCTACTACTGCCAGCTGGAGGCCAGT 1866
          ||| ||   ||| || || | | || |   || ||||||| |||| || || |
Db   1771 GTGATGTCAGTGGAAGATGAATCTACATC------CTGTTACTGCCTTTTGGACCCCTTT 1824

Qy   1867 GCCTGCTACGTCTTCACCGAGCAGCTGGGCCGCTTTGCCCTGGTGGGAGAGGCCCTCAGC 1926
          || || | || || || | ||    | ||  ||| || ||| ||      ||||||| |||
Db   1825 GCGTGTCATGTGCTCCTGGACAGCTTTGGGACCTATGCGCTCACTGGAGAGCCAATCACA 1884

Qy   1927 GTGGCTGCCGCCAAGCGCCCTCAAGCTGCTTCTGTTTGCGCCGGTGGCCTGCACCTCCCTC 1986
          |    |||||  ||||  || ||| ||  | ||||      || |||| | |||||||
Db   1885 GACTGTGCCGTGAAGCAACTGAAGGTGGCGGTTTTTGGCTGCATGTCCTGTAACTCCCTG 1944
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Qy    1987 GAGTACAACATCCGGGTCTACTGCCTGCATGACACCCACGATGCACTCAAGGAGGTGGTG 2046
           || |||||||  |  | || |||||  || |  | ||||   |||| |  |||| |||||
Db    1945 GATTACAACTTGAGAGTTTACTGTGTGGACAATACCCCTTGTGCATTTCAGGAAGTGGTT 2004

Qy    2047 CAGCTGGAGAAGCAGCTGGGGGGACAGCTGATCCAGGAGCCACGGGTCCTGCACTTCAAG 2106
           || | ||| |  || |||||||||  |  | || |||     | ||||| |||||
Db    2005 TCAGATGAAAGGCATCAAGGTGGACAGCTCCTGGAAGAACCAAAATTGCTGCATTTCAAA 2064

Qy    2107 GACAGTTACCACAACCTGCGCCTATCCATCCACGATGTGCCCAGCTCCCTGTGGAAGAGT 2166
           |  |  | |   |   |  || |  | ||   ||| | ||| | ||| |||| | |
Db    2065 GGGAATACCTTTAGTCTTCAGATTTCTGTCCTTGATATTCCCCCATTCCTCTGGAGAATT 2124

Qy    2167 AAGCTCCTTGTCAGCTACCAGGAGATCCCCTTTTATCACATCTGGAATGGCACGCAGCGG 2226
           || |   |      || |||||| |||| ||  |  | | |||    | |  | || |
Db    2125 AAACCATTCACTGCCTGCCAGGAAGTCCCGTTCTCCCGCGTGTGGTGCAGTAACCGGCAG 2184

Qy    2227 TACTTGCACTGCACCTTCACCCTGGAGCGTGTCAGCCCCAGCACTAGTGACCTGGCCTGC 2286
            | |||||||| |||||| ||||||||||||| |  | ||||  || |   | ||| |||||
Db    2185 CCCCTGCACTGTGCCTTCTCCCTGGAGCGTTATACGCCCACTACCACCCAGCTGTCCTGC 2244

Qy    2287 AAGCTGTGGGTGTGGCAGGTGGAGGGCGACGGGCAGAGCTTCAGCATCAACTTCAACATC 2346
           ||  |  || |  ||||| |  | ||| |  |||| | || |  | |        |||
Db    2245 AAAATCTGCATTCGGCAGCTCAAAGGCCATGAACAGATCCTCCAAGTGCAGACATCAATC 2304

Qy    2347 ACCAAGGACACAAGGTTTGCTGAGCTGCTGGCTCTGGAGAGTGAAGCGGGGGTCCCAGCC 2406
             ||     | |    |      | |    |    |  | ||    |||| ||
Db    2305 CTAGAGAGTGAACGAGAAACCATCACTTTCTTCGCACAAGAGGACAGCACTTTCCCTGCA 2364

Qy    2407 CTGGTGGGCCCCAGTGCCTTCAAGATCCCCTTCCTCATTCGGCAGAAGATAATTTCCAGC 2466
           | |     |||||||| |||||||||| || |||| |  ||| | |||  ||  | | |
Db    2365 CAGACTGGCCCCAAAGCCTTCAAAATTCCCTACTCCATCAGACAGCGGATTTGTGCTACA 2424

Qy    2467 CTGGACCCACCCTGTAGGCGGGGTGCCGACTGGCGGACTCTGGCCCAGAAACTCCACCTG 2526
           | || | ||| |   || | ||||||| || | || |||||| | | |
Db    2425 TTTGATACCCCCAATGCCAAAGGCAAGGACTGGCAGATGTTAGCACAGAAAAACAGCATC 2484

Qy    2527 GACAGCCATCTCAGCTTCTTTGCCTCCAAGCCCAGCCCCACAGCCATGATCCTCAACCTG 2586
           ||||  || |     |  || ||  | |     ||||| |  || |  | |||||||
Db    2485 AACAGGAATTTATCTTATTTCGCTACACAAAGTAGCCCATCTGCTGTCATTTTGAACCTG 2544

Qy    2587 TGGGAGGCGCGGCACTTCCCCAACGGCAACCTCAGCCAGCTGGCTGCAGCAGTGGCTGGA 2646
           ||||| || || || ||   |  | ||  | ||  |  |||||   ||  | | |
Db    2545 TGGGAAGCTCGTCATCAGCATGATGGTGATCTTGACTCCCTGGCCTGTGCCCTTGAAGAG 2604

Qy    2647 CTGGGCCAGCCAGACGCTGGCCTCTTCACAGTGTCGGAGGCTGAG 2691
           |  ||  | || || |  |||| |  | || || |  |  ||
Db    2605 ATTGGGAGGACACACACGAAACTCTCAAACATTTCAGAATCCCAG 2649
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RESULT 12
US-09-969-532-15
; Sequence 15, Application US/09969532
; Patent No. 6777232
; GENERAL INFORMATION:
;   APPLICANT: Walke, D. Wade
;   APPLICANT: Scoville, John

```
  Query Match              16.3%;   Score 439.8;   DB 4;   Length 2661;
  Best Local Similarity    51.2%;   Pred. No. 2.3e-90;
  Matches 1313;  Conservative    0;  Mismatches 1147;  Indels  105;  Gaps    8;


Qy       127 CTTCCCCACTTCCTGGTGGAGCCCGAGGATGTGTACATCGTCAAGAACAAGCCAGTGCTG 186
             || || || ||| | | |||||| || |||| || ||  |||||| ||| || |
Db       157 CTGCCTCATTTCATAGAGGAGCCAGATGATGCTTATATTATCAAGAGCAACCCTATTGCA 216


Qy       187 CTTGTGTGCAAGGCCGTGCCCGCCACGCAGATCTTCTTCAAGTGCAACGGGGAGTGGGTG 246
             ||    ||||||| ||    ||| |||| |||||| |||||||||| |||||||| ||||||||
Db       217 CTCAGGTGCAAAGCGAGGCCAGCCATGCAGATATTCTTCAAATGCAACGGCGAGTGGGTC 276


Qy       247 CGCCAGGTGGACCACGTGATCGAGCGCAGCACAGACGGGAGCAGTGGGCTGCCCACCATG 306
             |  |||   || |||||     ||   |      |||| ||||  || ||       |
Db       277 CATCAGAACGAGCACGTCTCTGAAGAGACTCTGGACGAGAGCTCAGGTTTGAAGGTCCGC 336


Qy       307 GAGGTCCGCATTAATGTCTCAAGGCAGCAGGTCGAGAAGGTGTTCGGGCTGGAGGAATAC 366
             || ||    ||| |||||  | |||||| |||||| ||| | |     |||| ||||| ||
Db       337 GAAGTGTTCATCAATGTTACTAGGCAACAGGTGGAGGACTTCCATGGGCCCGAGGACTAT 396


Qy       367 TGGTGCCAGTGCGTGGCATGGAGCTCCTCGGGCACCACCAAGAGTCAGAAGGCCTACATC 426
             |||||||||||| ||||| |||||| |   ||| ||| ||||||||    |||||||| |
Db       397 TGGTGCCAGTGTGTGGCGTGGAGCCACCTGGGTACCTCCAAGAGCAGGAAGGCCTCTGTG 456


Qy       427 CGCATAGCCAGATTGCGCAAGAACTTCGAGCAGGAGCCGCTGGCCAAGGAGGTGTCCCTG 486
             |||||||||    || || || ||||| || || || || || |  |  | | ||| || | |
Db       457 CGCATAGCCTATTTACGGAAAAACTTTGAACAAGACCCACAAGGAAGGGAAGTTCCCATT 516


Qy       487 GAGCAGGGCATCGTGCTGCCCTGCCGTCCACCGGAGGGCATCCCTCCAGCCGAGGTGGAG 546
             ||       || || |||| ||||||| ||||| ||||| |||||  ||||| | ||||||||||||
Db       517 GAAGGCATGATTGTACTGCACTGCCGCCCACCAGAGGGAGTCCCTGCTGCCGAGGTGGAA 576


Qy       547 TGGCTCCGGAACGAGGACCTGGTGGACCCGTCCCTGGACCCCAATGTATACATCACGCGG 606
             |||||    || || || ||  | ||| |    |  ||| || |   ||| || |
Db       577 TGGCTGAAAAATGAAGAGCCCATTGACTCTGAACAAGACGAGAACATTGACACCAGGGCT 636


Qy       607 GAGCACAGCCTGGTGGTGCGACAGGCCCGCCTTGCTGACACGGCCAACTACACCTGCGTG 666
             || || | |||| |   | | ||||| || ||   | ||| | |  || ||||||||||| ||
Db       637 GACCATAACCTGATCATCAGGCAGGCACGGCTCTCGGACTCAGGAAATTACACCTGCATG 696


Qy       667 GCCAAGAACATCGTGGCACGTCGCCGCAGCGCCTCCGCTGCTGTCATCGTCTACGTGAAC 726
             ||     |||||||||||         |  | |||    || ||  |||| | |||||||||| |
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Db    697 GCAGCCAACATCGTGGCTAAGAGGAGAAGCCTGTCGGCCACTGTTGTGGTCTACGTGGAT 756

Qy    727 GGTGGGTGGTCGACGTGGACCGAGTGGTCCGTCTGCAGCGCCAGCTGTGGGCGCGGCTGG 786
              ||  | |||      |||||  |||  |||||||||||||||||  |    ||||
Db    757 GGGAGCTGGGAAGTGTGGAGCGAATGGTCCGTCTGCAGTCCAGAGTGTG--------AA 807

Qy    787 CAGAAACGGAGCCGGAGCTGCACCAACCCGGCGCCTCTCAACGGGGGCGCTTTCTGTGAG 846
              ||    |||| ||||     |||||    || | ||    || ||||||   ||||||||
Db    808 CATTTGCGGATCCGGGAGTGCACAGCACCACCCCCGAGAAATGGGGGCAAATTCTGTGAA 867

Qy    847 GGGCAGAATGTCCAGAAAACAGCCTGCGCCACCCTGTGCCCAGTAGACGGCAGCTGGAGC 906
              || |  |        |       |  |||| |             |    ||||    ||
Db    868 GGTCTAAGCCAGGAATCTGAAAACTGCACAGATGGTCTTTGCATCCTAGGCATTGAGAAT 927

Qy    907 CCGTGGAGCAAGTGGTCGGCCTGTGGGCTGGACTGCACCCACTGGCGGAGCCGTGAGTGC 966
               |  |   ||   |   || || ||  ||  |||
Db    928 GCCAGCGACATTGCTTTGTACTCGGGCTTGG----------------------------- 958

Qy    967 TCTGACCCAGCACCCCGCAACGGAGGGGAGGAGTGCCAGGGCACTGACCTGGACACCCGC 1026
                      |     |   | | |   ||     ||  ||    | | | |      |  |
Db    959 ---------------GTGCTGCCGTCGTGGCCGTTGCAGTCCTGGTCATTGGTGTCACC 1002

Qy   1027 AACTGTACCAGTGACCTCTGTGTACACAGTGCTTCTGGCCCTGAGGACGTGGCCCTCTAT 1086
              |   |    |   ||    ||| |   |   || |    ||  ||||||      ||  |
Db   1003 CTTTACAGACGGAGCCAGAGTGACTATGGCGTGGACGTCATTGACTCTTCTGCATTGACA 1062

Qy   1087 GTGGGCCTCATCGCCGTGGCCGTCTGCCTGGTCCTGCTGCTGCTTGTCCTCATCCTCGTT 1146
              | |||  ||      || |   | ||    |||| |        ||| ||||   |
Db   1063 GGTGGCTTCCAGACCTTCAACTTCAAAACAGTCCGTCAAGGTAACTCCCTGCTCCTGAAT 1122

Qy   1147 TATTGCCGGAAGAAGGAGGGGCTGGACTCAGATGTGGCTGACTCGTCCATTCTCACCTCA 1206
              | |  | |  ||      | | ||  || ||  |  ||  |      |    | | | ||
Db   1123 TCTGCCATGCA--------GCCAGATCTGACAGTGAGCCGGACATACAGCGGACCCATCT 1174

Qy   1207 GGCTTCCAGCCCGTCAGCATCAAGCCCAGCAAAGCAGACAACCCCCATCTGCTCACCATC 1266
              | || | | || || | |  | | | |   ||| | | | |  || ||     | ||| |
Db   1175 GTCTGCAGGACCCTCTGGACAAGG---AGCTCATGACAGAGTCCTCACTCTTTAACCCTT 1231

Qy   1267 CAGCCGGACCTCAGCACCACCACCACCACCTACCAGGGCAGTCTCTGTCCCCGGCAGGAT 1326
               | |        |   | |    ||| |      |   | || |||    | ||
Db   1232 TGTCGGACATCAAAGTGAAAGTCCAGAGCTCGTTCATGGTTTCCCTGGGAGTGTCTGAGA 1291

Qy   1327 GGGCCCAGCCCCAAGTTCCAGCTCACCAATGGGCACCTGCTCAGCCCCCTGGGTGGCGGC 1386
              | ||  ||  ||| |       ||| |       || | |
Db   1292 GAGCTGAGTACCACGGCAAGAATCATTCCAGGACTTTT---------------------- 1329

Qy   1387 CGCCACACACTGCACCACAGCTCTCCCACCTCTGAGGCCGAGGAGTTCGTCTCCCGCCTC 1446
              | |||    |  ||||||||||| | ||    ||   | || |    |   |  || | ||
Db   1330 CCCCATGGAAACAACCACAGCTTTAGTACAATGCATCCCAGAAATAAAATGCCCTACATC 1389

Qy   1447 TCCACCCAGAACTACTTCCGCTCCCTGCCCCGAGGCACCAGCAACATGACCTATGGGACC 1506
              |   | |       || || |||  | | | ||       | |||     ||| |   |
Db   1390 CAAAATCTGT---------CATCACTCCCCACAAGGACAGAACTGAGGACAACTGGTGTC 1440

Qy   1507 TTCAACTTCCTCGGGGGCCGGCTGATGATCCCTAATACAGGTATCAGCCTCCTCATCCCC 1566
              ||   |    | |||||  ||  | | || || ||||||||| | ||| | |||||| ||
Db   1441 TTTGGCCATTTAGGGGGGCGCTTAGTAATGCCAAATACAGGGGTGAGCTTACTCATACCA 1500
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Qy    1567 CCAGATGCCATACCCCGAGGGAAGATCTATGAGATCTACCTCACGCTGCACAAGCCGGAA.1626
           | | |||||| ||    | |||     |  ||||| ||  | | |    ||| | |
Db    1501 CACGGTGCCATCCCAGAGGAGAATTCTTGGGAGATTTATATGTCCAT---CAACCAAGGT 1557

Qy    1627 GACGTGAGGTTGCCCCTAGCTGGCTGTCAGACCCTGCTGAGTCCCATCGTTAGCTGTGGA 1686
           ||     ||  | |   ||  |||||| | ||  || |||||||||    || | ||||||
Db    1558 GAACCCAGCCTCCAGTCAGATGGCTCTGAGGTGCTCCTGAGTCCTGAAGTCACCTGTGGT 1617

Qy    1687 CCCCCTGGCGTCCTGCTCACCCGGCCAGTCATCCTGGCTATGGACCACTGTGGGGAGCCC 1746
           || || | | |   |||||  || ||  | |   || | ||  |||||||| ||    |
Db    1618 CCTCCAGACATGATCGTCACCACTCCCTTTGCATTGACCATCCCGCACTGTGCAGATGTC 1677

Qy    1747 AGCCCTGACAGCTGGAGCCTGCGCCTCAAAAAGCAGTCGTGCGAGGGCAGCTGGGAGGAT 1806
           ||  ||||    ||||    | |  | || ||| | |   |||||| ||||||||||||||
Db    1678 AGTTCTGAGCATTGGAATATCCATTTAAAGAAGAGGACACAGCAGGGCAAATGGGAGGAA 1737

Qy    1807 GTGCTGCACCTGGGCGAGGAGGCGCCCTCCCACCTCTACTACTGCCAGCTGGAGGCCAGT 1866
           ||| ||    ||| || || || | | ||    || ||||||| ||||   || ||  |
Db    1738 GTGATGTCAGTGGAAGATGAATCTACATC------CTGTTACTGCCTTTTGGACCCCTTT 1791

Qy    1867 GCCTGCTACGTCTTCACCGAGCAGCTGGGCCGCTTTGCCCTGGTGGGAGAGGCCCTCAGC 1926
           || ||  | || ||   || | | ||   | ||   || ||||||| |||| |  |||
Db    1792 GCGTGTCATGTGCTCCTGGACAGCTTTGGGACCTATGCGCTCACTGGAGAGCCAATCACA 1851

Qy    1927 GTGGCTGCCGCCAAGCGCCTCAAGCTGCTTCTGTTTGCGCCGGTGGCCTGCACCTCCCTC 1986
           |     ||||| ||||  | |||| || || |  | ||||   || |||| | |||||||
Db    1852 GACTGTGCCGTGAAGCAACTGAAGGTGGCGGTTTTTGGCTGCATGTCCTGTAACTCCCTG 1911

Qy    1987 GAGTACAACATCCGGGTCTACTGCCTGCATGACACCCACGATGCACTCAAGGAGGTGGTG 2046
           || |||||| |  | || ||||| ||  | | | |||| |||| |  |||| ||| |||||
Db    1912 GATTACAACTTGAGAGTTTACTGTGTGGACAATACCCCTTGTGCATTTCAGGAAGTGGTT 1971

Qy    2047 CAGCTGGAGAAGCAGCTGGGGGGACAGCTGATCCAGGAGCCACGGGTCCTGCACTTCAAG 2106
           || | ||| |  || |||||||| |   | || |||    | ||||| |||||
Db    1972 TCAGATGAAAGGCATCAAGGTGGACAGCTCCTGGAAGAACCAAAATTGCTGCATTTCAAA 2031

Qy    2107 GACAGTTACCACAACCTGCGCCTATCCATCCACGATGTGCCCAGCTCCCTGTGGAAGAGT 2166
           |  |  |    |   |  || |  | ||  ||| ||| | |  ||  | ||| |||| | |
Db    2032 GGGAATACCTTTAGTCTTCAGATTTCTGTCCTTGATATTCCCCCATTCCTCTGGAGAATT 2091

Qy    2167 AAGCTCCTTGTCAGCTACCAGGAGATCCCCTTTTATCACATCTGGAATGGCACGCAGCGG 2226
           || |   |    ||  ||||||  |||||| | ||| | | ||    | |  ||| | |
Db    2092 AAACCATTCACTGCCTGCCAGGAAGTCCCGTTCTCCCGCGTGTGGTGCAGTAACCGGCAG 2151

Qy    2227 TACTTGCACTGCACCTTCACCCTGGAGCGTGTCAGCCCCAGCACTAGTGACCTGGCCTGC 2286
           |  |||||||  |||||  ||||| ||||||| ||     | |||| || |  ||| |||||
Db    2152 CCCCTGCACTGTGCCTTCTCCCTGGAGCGTTATACGCCCACTACCACCCAGCTGTCCTGC 2211

Qy    2287 AAGCTGTGGGTGTGGCAGGTGGAGGGCGACGGGCAGAGCTTCAGCATCAACTTCAACATC 2346
           ||  |  | ||  | |||||| |  | ||| | |  |||| | ||    | |      |||
Db    2212 AAAATCTGCATTCGGCAGCTCAAAGGCCATGAACAGATCCTCCAAGTGCAGACATCAATC 2271

Qy    2347 ACCAAGGACACAAGGTTTGCTGAGCTGCTGGCTCTGGAGAGTGAAGCGGGGGTCCCAGCC 2406
           ||     || |  |    |   |  |     | |||    |  || ||    |||| ||
Db    2272 CTAGAGAGTGAACGAGAAACCATCACTTTCTTCGCACAAGAGGACAGCACTTTCCCTGCA 2331
```

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Qy    2407  CTGGTGGGCCCCAGTGCCTTCAAGATCCCCTTCCTCATTCGGCAGAAGATAATTTCCAGC  2466
             | |   | |||||||   ||||||||  ||  ||||  |   ||| | |||  |||   | | |
Db    2332  CAGACTGGCCCCAAAGCCTTCAAAATTCCCTACTCCATCAGACAGCGGATTTGTGCTACA  2391

Qy    2467  CTGGACCCACCCTGTAGGCGGGGTGCCGACTGGCGGACTCTGGCCCAGAAACTCCACCTG  2526
             | ||  | |||  |       ||     |||||||| ||  | ||  ||||||| |   | |
Db    2392  TTTGATACCCCCAATGCCAAAGGCAAGGACTGGCAGATGTTAGCACAGAAAAACAGCATC  2451

Qy    2527  GACAGCCATCTCAGCTTCTTTGCCTCCAAGCCCAGCCCCACAGCCATGATCCTCAACCTG  2586
             ||||  || |    |  || ||  | |     |||||  | ||  | ||  | ||||||
Db    2452  AACAGGAATTTATCTTATTTCGCTACACAAAGTAGCCCATCTGCTGTCATTTTGAACCTG  2511

Qy    2587  TGGGAGGCGCGGCACTTCCCCAACGGCAACCTCAGCCAGCTGGCTGCAGCAGTGGCTGGA  2646
             |||||  || ||  ||      |   | ||  | ||     |  |||||    ||  | | |
Db    2512  TGGGAAGCTCGTCATCAGCATGATGGTGATCTTGACTCCCTGGCCTGTGCCCTTGAAGAG  2571

Qy    2647  CTGGGCCAGCCAGACGCTGGCCTCTTCACAGTGTCGGAGGCTGAG  2691
             | ||   | || || |    ||||  |   | || ||  |  ||
Db    2572  ATTGGGAGGACACACACGAAACTCTCAAACATTTCAGAATCCCAG  2616
```

RESULT 13
US-09-471-276-345
; Sequence 345, Application US/09471276
; Patent No. 6822072
; GENERAL INFORMATION:
; APPLICANT: Dumas Milne Edwards, J.B.
; APPLICANT: Duclert A.
; APPLICANT: Giordano, J.Y.
; TITLE OF INVENTION: Expressed Sequence Tags and Encoded Human Proteins.
; Patent No. 6822072
; FILE REFERENCE: GENSET.025CP1
; CURRENT APPLICATION NUMBER: US/09/471,276
; CURRENT FILING DATE: 1999-12-21
; EARLIER APPLICATION NUMBER: 09/057,719
; EARLIER FILING DATE: 1998-04-09
; EARLIER APPLICATION NUMBER: 09/069,047
; EARLIER FILING DATE: 1998-04-28
; EARLIER APPLICATION NUMBER: PCT/IB99/00712
; EARLIER FILING DATE: 1999-04-09
; NUMBER OF SEQ ID NOS: 1622
; SOFTWARE: Patent.pm
; SEQ ID NO 345
; LENGTH: 349
; TYPE: DNA
; ORGANISM: Homo sapiens
; FEATURE:
; NAME/KEY: CDS
; LOCATION: 207..347
; NAME/KEY: sig_peptide
; LOCATION: 207..278
; OTHER INFORMATION: Von Heijne matrix
; OTHER INFORMATION: score 5.40000009536743
; OTHER INFORMATION: seq SCCCLSSSSFIAG/RR
US-09-471-276-345

   Query Match   12.0%; Score 323; DB 4; Length 349;

Best Local Similarity   98.8%;   Pred. No. 4.4e-64;
Matches   335;   Conservative   1;   Mismatches   2;   Indels   1;   Gaps   1;

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Qy       889 GTAGACGGCAGCTGGAGCCCGTGGAGCAAGTGGTCGGCCTGTGGGCTGGACTGCACCCAC 948
             || ||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        12 GTGGACGGCAGCTGGAGCCCGTGGAGCAAGTGGTCGGCCTGTGGGCTGGACTGCACCCAC 71

Qy       949 TGGCGGAGCCGTGAGTGCTCTGACCCAGCACCCCGCAACGGAGGGGAGGAGTGCCAGGGC 1008
             ||||||| ||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        72 TGGCGGA-CCGTGAGTGCTCTGACCCAGCACCCCGCAACGGAGGGGAGGAGTGCCAGGGC 130

Qy      1009 ACTGACCTGGACACCCGCAACTGTACCAGTGACCTCTGTGTACACAGTGCTTCTGGCCCT 1068
             ||||||||||||||||||||||||||||||||||||||||||||||| ||||||||||||
Db       131 ACTGACCTGGACACCCGCAACTGTACCAGTGACCTCTGTGTACACACTGCTTCTGGCCCT 190

Qy      1069 GAGGACGTGGCCCTCTATGTGGGCCTCATCGCCGTGGCCGTCTGCCTGGTCCTGCTGCTG 1128
             |||||||||||||||||||||||||||||||||||||||||||||:|||||||||||||
Db       191 GAGGACGTGGCCCTCTATGTGGGCCTCATCGCCGTGGCCGTCTGCBTGGTCCTGCTGCTG 250

Qy      1129 CTTGTCCTCATCCTCGTTTATTGCCGGAAGAAGGAGGGGCTGGACTCAGATGTGGCTGAC 1188
             ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db       251 CTTGTCCTCATCCTCGTTTATTGCCGGAAGAAGGAGGGGCTGGACTCAGATGTGGCTGAC 310

Qy      1189 TCGTCCATTCTCACCTCAGGCTTCCAGCCCGTCAGCATC 1227
             |||||||||||||||||||||||||||||||||||||||
Db       311 TCGTCCATTCTCACCTCAGGCTTCCAGCCCGTCAGCATC 349
```

RESULT 14
US-09-969-532-31
; Sequence 31, Application US/09969532
; Patent No. 6777232
; GENERAL INFORMATION:
;   APPLICANT: Walke, D. Wade
;   APPLICANT: Scoville, John
;   TITLE OF INVENTION: No. 6777232el Human Membrane Proteins and Polynucleotides
Encoding the Same
;   FILE REFERENCE: LEX-0244-USA
;   CURRENT APPLICATION NUMBER: US/09/969,532
;   CURRENT FILING DATE: 2001-10-02
;   PRIOR APPLICATION NUMBER: US 60/237,280
;   PRIOR FILING DATE: 2000-10-02
;   NUMBER OF SEQ ID NOS: 33
;   SOFTWARE: FastSEQ for Windows Version 4.0
; SEQ ID NO 31
;   LENGTH: 1968 .
;   TYPE: DNA
;   ORGANISM: homo sapiens
US-09-969-532-31

Query Match   10.9%;   Score 293.2;   DB 4;   Length 1968;
Best Local Similarity   53.7%;   Pred. No. 4.9e-57;
Matches   657;   Conservative   0;   Mismatches   558;   Indels   9;   Gaps   2;

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Qy      1468 TCCCTGCCCCGAGGCACCAGCAACATGACCTATGGGACCTTCAACTTCCTCGGGGGCCGG 1527
             || || |||  | | ||        | ||| |||   ||| |   |  | |||||| ||
Db       709 TCACTCCCCACAAGGACAGAACTGAGGACAACTGGTGTCTTTGGCCATTTAGGGGGGCGC 768
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Qy     1528 CTGATGATCCCTAATACAGGTATCAGCCTCCTCATCCCCCCAGATGCCATACCCCGAGGG 1587
            |  |  || || |||||||||  |  |||  |  ||||| || |  | |||||| ||    | |
Db      769 TTAGTAATGCCAAATACAGGGGTGAGCTTACTCATACCACACGGTGCCATCCCAGAGGAG 828

Qy     1588 AAGATCTATGAGATCTACCTCACGCTGCACAAGCCGGAAGACGTGAGGTTGCCCCTAGCT 1647
            ||    |  ||||| ||  |  |  |  | ||  |    |||  |  || || ||     || |
Db      829 AATTCTTGGGAGATTTATATGTCCATCAACCAAGGTGAACCC---AGCCTCCAGTCAGAT 885

Qy     1648 GGCTGTCAGACCCTGCTGAGTCCCATCGTTAGCTGTGGACCCCCTGGCGTCCTGCTCACC 1707
            |||| | ||   || |||||||||     ||  | |||||||| || || | | |   ||||| 
Db      886 GGCTCTGAGGTGCTCCTGAGTCCTGAAGTCACCTGTGGTCCTCCAGACATGATCGTCACC 945

Qy     1708 CGGCCAGTCATCCTGGCTATGGACCACTGTGGGGAGCCCAGCCCTGACAGCTGGAGCCTG 1767
            ||  |      || | ||  |||||||  ||  || |||  ||||      ||||    |
Db      946 ACTCCCTTTGCATTGACCATCCCGCACTGTGCAGATGTCAGTTCTGAGCATTGGAATATC 1005

Qy     1768 CGCCTCAAAAAGCAGTCGTGCGAGGGCAGCTGGGAGGATGTGCTGCACCTGGGCGAGGAG 1827
            |   |  | || |||   | |  ||||||  ||||||||| ||| ||    ||| || || ||
Db     1006 CATTTAAAGAAGAGGACACAGCAGGGCAAATGGGAGGAAGTGATGTCAGTGGAAGATGAA 1065

Qy     1828 GCGCCCTCCCACCTCTACTACTGCCAGCTGGAGGCCAGTGCCTGCTACGTCTTCACCGAG 1887
            |   |  ||     || ||||||| |||| ||   || ||   |||| ||  || ||   ||
Db     1066 TCTACATC------CTGTTACTGCCTTTTGGACCCCTTTGCGTGTCATGTGCTCCTGGAC 1119

Qy     1888 CAGCTGGGCCGCTTTGCCCTGGTGGGAGAGGCCCTCAGCGTGGCTGCCGCCAAGCGCCTC 1947
            | ||   || ||| ||   ||  |||||| |   ||| |     ||||| ||||    ||
Db     1120 AGCTTTGGGACCTATGCGCTCACTGGAGAGCCAATCACAGACTGTGCCGTGAAGCAACTG 1179

Qy     1948 AAGCTGCTTCTGTTTGCGCCGGTGGCCTGCACCTCCCTCGAGTACAACATCCGGGTCTAC 2007
            ||| || | |||   || ||| || ||  ||||| |||||| | |||||| | || |||
Db     1180 AAGGTGGCGGTTTTTGGCTGCATGTCCTGTAACTCCCTGGATTACAACTTGAGAGTTTAC 1239

Qy     2008 TGCCTGCATGACACCCACGATGCACTCAAGGAGGTGGTGCAGCTGGAGAAGCAGCTGGGG 2067
            || || |  |  | |||| |||| | |||| | |||| |||| |      || |  ||| |  ||
Db     1240 TGTGTGGACAATACCCCTTGTGCATTTCAGGAAGTGGTTTCAGATGAAAGGCATCAAGGT 1299

Qy     2068 GGACAGCTGATCCAGGAGCCACGGGTCCTGCACTTCAAGGACAGTTACCACAACCTGCGC 2127
            ||||||| |  |  | || |||    | |||||| ||||| | | | |  |   |   || |
Db     1300 GGACAGCTCCTGGAAGAACCAAAATTGCTGCATTTCAAAGGGAATACCTTTAGTCTTCAG 1359

Qy     2128 CTATCCATCCACGATGTGCCCAGCTCCCTGTGGAAGAGTAAGCTCCTTGTCAGCTACCAG 2187
            | ||   |||   ||| | |||  | |||| ||||  | |||| |   |     || ||||
Db     1360 ATTTCTGTCCTTGATATTCCCCCATTCCTCTGGAGAATTAAACCATTCACTGCCTGCCAG 1419

Qy     2188 GAGATCCCCTTTTATCACATCTGGAATGGCACGCAGCGGTACTTGCACTGCACCTTCACC 2247..
            ||  |||| || |  | | | | |||  | |  | || |  | ||||||| |||||  ||
Db     1420 GAAGTCCCGTTCTCCCGCGTGTGGTGCAGTAACCGGCAGCCCCTGCACTGTGCCTTCTCC 1479

Qy     2248 CTGGAGCGTGTCAGCCCCAGCACTAGTGACCTGGCCTGCAAGCTGTGGGTGTGGCAGGTG 2307
            ||||||||| |    |||| || | ||  |  ||| ||||||||||| | || | ||||| |
Db     1480 CTGGAGCGTTATACGCCCACTACCACCCAGCTGTCCTGCAAAATCTGCATTCGGCAGCTC 1539

Qy     2308 GAGGGCGACGGGCAGAGCTTCAGCATCAACTTCAACATCACCAAGGACACAAGGTTTGCT 2367
            || |||| |   |||| | ||  | ||     | |    |||   || |   | |  |  |
Db     1540 AAAGGCCATGAACAGATCCTCCAAGTGCAGACATCAATCCTAGAGAGTGAACGAGAAACC 1599
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Qy     2368 GAGCTGCTGGCTCTGGAGAGTGAAGCGGGGGTCCCAGCCCTGGTGGGCCCCAGTGCCTTC 2427
                   |   .       |     | ||        |||| || | |    ||||||| ||||||
Db     1600 ATCACTTTCTTCGCACAAGAGGACAGCACTTTCCCTGCACAGACTGGCCCCAAAGCCTTC 1659

Qy     2428 AAGATCCCCTTCCTCATTCGGCAGAAGATAATTTCCAGCCTGGACCCACCCTGTAGGCGG 2487
                 || || |||| |   ||| | ||| ||| | | |   |  | ||  | |||  |
Db     1660 AAAATTCCCTACTCCATCAGACAGCGGATTTGTGCTACATTTGATACCCCCAATGCCAAA 1719

Qy     2488 GGTGCCGACTGGCGGACTCTGGCCCAGAAACTCCACCTGGACAGCCATCTCAGCTTCTTT 2547
                 ||      ||||||| ||   | || |||||| | | |  |||| ||| |    | ||
Db     1720 GGCAAGGACTGGCAGATGTTAGCACAGAAAAACAGCATCAACAGGAATTTATCTTATTTC 1779

Qy     2548 GCCTCCAAGCCCAGCCCCACAGCCATGATCCTCAACCTGTGGGAGGCGCGGCACTTCCCC 2607
                 || | |     |||||  | ||  | ||  | |||||||||||| || || ||      |
Db     1780 GCTACACAAAGTAGCCCATCTGCTGTCATTTTGAACCTGTGGGAAGCTCGTCATCAGCAT 1839

Qy     2608 AACGGCAACCTCAGCCAGCTGGCTGCAGCAGTGGCTGGACTGGGCCAGCCAGACGCTGGC 2667
                 | || | ||  |   ||||| ||  | | |   | ||  | || || |
Db     1840 GATGGTGATCTTGACTCCCTGGCCTGTGCCCTTGAAGAGATTGGGAGGACACACACGAAA 1899

Qy     2668 CTCTTCACAGTGTCGGAGGCTGAG 2691
                 ||||  |   | || ||   | ||
Db     1900 CTCTCAAACATTTCAGAATCCCAG 1923


RESULT 15
US-09-969-532-29
; Sequence 29, Application US/09969532
; Patent No. 6777232
; GENERAL INFORMATION:
;  APPLICANT: Walke, D. Wade
;  APPLICANT:  Scoville, John
;   TITLE OF INVENTION: No. 6777232el Human Membrane Proteins and Polynucleotides
Encoding the Same
;  FILE REFERENCE: LEX-0244-USA
;  CURRENT APPLICATION NUMBER: US/09/969,532
;  CURRENT FILING DATE:  2001-10-02
;  PRIOR APPLICATION NUMBER: US 60/237,280
;  PRIOR FILING DATE: 2000-10-02
;  NUMBER OF SEQ ID NOS: 33
;  SOFTWARE: FastSEQ for Windows Version 4.0
; SEQ ID NO 29
;   LENGTH: 2001
;   TYPE: DNA
;   ORGANISM: homo sapiens
US-09-969-532-29

  Query Match            10.9%;   Score 293.2;  DB 4;  Length 2001;
  Best Local Similarity   53.7%;   Pred. No. 4.9e-57;
  Matches  657;  Conservative    0;  Mismatches  558;  Indels    9;  Gaps    2;

Qy     1468 TCCCTGCCCCGAGGCACCAGCAACATGACCTATGGGACCTTCAACTTCCTCGGGGGCCGG 1527
                 || || |||  | | ||        | |||   |||   |||   |    | |||||| ||
Db      742 TCACTCCCCACAAGGACAGAACTGAGGACAACTGGTGTCTTTGGCCATTTAGGGGGGCGC 801

Qy     1528 CTGATGATCCCTAATACAGGTATCAGCCTCCTCATCCCCCCAGATGCCATACCCCGAGGG 1587
                 |  | || || ||||||||| | ||| | ||||| || | | ||||||| ||     | |
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Db    802 TTAGTAATGCCAAATACAGGGGTGAGCTTACTCATACCACACGGTGCCATCCCAGAGGAG 861

Qy   1588 AAGATCTATGAGATCTACCTCACGCTGCACAAGCCGGAAGACGTGAGGTTGCCCCTAGCT 1647
           ||    |  ||||| ||   |  |   || |    ||| |   || | |    || |
Db    862 AATTCTTGGGAGATTTATATGTCCATCAACCAAGGTGAACCC---AGCCTCCAGTCAGAT 918

Qy   1648 GGCTGTCAGACCCTGCTGAGTCCCATCGTTAGCTGTGGACCCCCTGGCGTCCTGCTCACC 1707
           |||| |  ||    || ||||||||    || | |||||| || || | |  |   |||||
Db    919 GGCTCTGAGGTGCTCCTGAGTCCTGAAGTCACCTGTGGTCCTCCAGACATGATCGTCACC 978

Qy   1708 CGGCCAGTCATCCTGGCTATGGACCACTGTGGGGAGCCCAGCCCTGACAGCTGGAGCCTG 1767
            ||   |     || | ||  |||||||  ||   |||  ||||      ||||    |
Db    979 ACTCCCTTTGCATTGACCATCCCGCACTGTGCAGATGTCAGTTCTGAGCATTGGAATATC 1038

Qy   1768 CGCCTCAAAAAGCAGTCGTGCGAGGGCAGCTGGGAGGATGTGCTGCACCTGGGCGAGGAG 1827
           |  | || |||   ||  ||||||  |||||||| ||| ||    |||   || || ||
Db   1039 CATTTAAAGAAGAGGACACAGCAGGGCAAATGGGAGGAAGTGATGTCAGTGGAAGATGAA 1098

Qy   1828 GCGCCCTCCCACCTCTACTACTGCCAGCTGGAGGCCAGTGCCTGCTACGTCTTCACCGAG 1887
           |   | ||     || ||||||| |||| ||  || || ||| || | || || ||
Db   1099 TCTACATC------CTGTTACTGCCTTTTGGACCCCTTTGCGTGTCATGTGCTCCTGGAC 1152

Qy   1888 CAGCTGGGCCGCTTTGCCCTGGTGGGAGAGGCCCTCAGCGTGGCTGCCGCCAAGCGCCTC 1947
            | ||    || ||| ||   |||||| |   ||| |    ||||| |||||    ||
Db   1153 AGCTTTGGGACCTATGCGCTCACTGGAGAGCCAATCACAGACTGTGCCGTGAAGCAACTG 1212

Qy   1948 AAGCTGCTTCTGTTTGCGCCGGTGGCCTGCACCTCCCTCGAGTACAACATCCGGGTCTAC 2007
           ||| ||     | ||||     || |||| | |||||| || |||||| |  | || |||
Db   1213 AAGGTGGCGGTTTTTGGCTGCATGTCCTGTAACTCCCTGGATTACAACTTGAGAGTTTAC 1272

Qy   2008 TGCCTGCATGACACCCACGATGCACTCAAGGAGGTGGTGCAGCTGGAGAAGCAGCTGGGG 2067
           ||   || |  |  | ||||   |||| | |||| |||||   || | ||| | || ||
Db   1273 TGTGTGGACAATACCCCTTGTGCATTTCAGGAAGTGGTTTCAGATGAAAGGCATCAAGGT 1332

Qy   2068 GGACAGCTGATCCAGGAGCCACGGGTCCTGCACTTCAAGGACAGTTACCACAACCTGCGC 2127
           |||||||  |   | || |||    | |||||| ||||| | | | |  | | || | |
Db   1333 GGACAGCTCCTGGAAGAACCAAAATTGCTGCATTTCAAAGGGAATACCTTTAGTCTTCAG 1392

Qy   2128 CTATCCATCCACGATGTGCCCAGCTCCCTGTGGAAGAGTAAGCTCCTTGTCAGCTACCAG 2187
            | ||   |||   ||| | |||   | ||| |||| | |  |  |  |   || ||||
Db   1393 ATTTCTGTCCTTGATATTCCCCCATTCCTCTGGAGAATTAAACCATTCACTGCCTGCCAG 1452

Qy   2188 GAGATCCCCTTTTATCACATCTGGAATGGCACGCAGCGGTACTTGCACTGCACCTTCACC 2247
           ||  ||||· ||  |  | | |||   |  |  || | ||||||| |||||| ||
Db   1453 GAAGTCCCGTTCTCCCGCGTGTGGTGCAGTAACCGGCAGCCCCTGCACTGTGCCTTCTCC 1512

Qy   2248 CTGGAGCGTGTCAGCCCCAGCACTAGTGACCTGGCCTGCAAGCTGTGGGTGTGGCAGGTG 2307
           ||||||||    |   |||| || | |   |||   | ||||||||| | || | ||||| |
Db   1513 CTGGAGCGTTATACGCCCACTACCACCCAGCTGTCCTGCAAAATCTGCATTCGGCAGCTC 1572

Qy   2308 GAGGGCGACGGGCAGAGCTTCAGCATCAACTTCAACATCACCAAGGACACAAGGTTTGCT 2367
           | ||| | |  |||| | ||   |    |  |||   ||    || |  |    |
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Search completed: March  6, 2005, 10:24:58
Job time : 441.67 secs
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GenCore version 5.1.6
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OM nucleic - nucleic search, using sw model

Run on: March 6, 2005, 05:25:16 ; Search time 1458.63 Seconds
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Scoring table: IDENTITY_NUC
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Searched: 5401638 seqs, 2966923429 residues

Total number of hits satisfying chosen parameters: 10803276

Minimum DB seq length: 0
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Post-processing: Minimum Match 0%
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        Pred. No. is the number of results predicted by chance to have a
        score greater than or equal to the score of the result being printed,
        and is derived by analysis of the total score distribution.

Result No.	Score	% Query Match	Length	DB	ID	Description
1	2697	100.0	2752	10	US-09-918-779-1	Sequence 1, Appli
2	2697	100.0	2752	17	US-10-624-932-1	Sequence 1, Appli
3	2621.4	97.2	2881	10	US-09-970-944-1	Sequence 1, Appli
4	2343	86.9	3561	18	US-10-643-795A-77	Sequence 77, Appl
5	2343	86.9	3580	17	US-10-311-623-13	Sequence 13, Appl
6	2252.2	83.5	2697	16	US-10-240-154-15	Sequence 15, Appl
7	2252.2	83.5	3014	10	US-09-933-261-1	Sequence 1, Appli
8	2252.2	83.5	3014	14	US-10-256-702-1	Sequence 1, Appli
9	1552.4	57.6	1787	10	US-09-933-261-2	Sequence 2, Appli
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11	1200.6	44.5	1321	17	US-10-296-115-365	Sequence 365, App
12	936.2	34.7	2860	17	US-10-087-684-1	Sequence 1, Appli
13	936.2	34.7	2860	17	US-10-087-684-3	Sequence 3, Appli
14	936.2	34.7	2860	17	US-10-218-779-1	Sequence 1, Appli
15	936.2	34.7	2860	17	US-10-218-779-3	Sequence 3, Appli
16	912.2	33.8	2895	17	US-10-037-417-37	Sequence 37, Appl
17	902.6	33.5	3485	9	US-09-816-828-18	Sequence 18, Appl
18	901	33.4	3884	14	US-10-028-072-145	Sequence 145, App
19	901	33.4	3884	14	US-10-140-808-145	Sequence 145, App
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23	901	33.4	3884	14	US-10-175-746-145	Sequence 145, App
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25	901	33.4	3884	14	US-10-176-921-145	Sequence 145, App
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27	901	33.4	3884	14	US-10-140-474-145	Sequence 145, App
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32	901	33.4	3884	14	US-10-142-423-145	Sequence 145, App
33	901	33.4	3884	14	US-10-121-050-145	Sequence 145, App
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37	901	33.4	3884	14	US-10-123-236-145	Sequence 145, App
38	901	33.4	3884	14	US-10-123-261-145	Sequence 145, App
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ALIGNMENTS

RESULT 1
US-09-918-779-1

; Sequence 1, Application US/09918779
; Publication No. US20030064369A1
; GENERAL INFORMATION:
; APPLICANT: Taupier, Raymond
; APPLICANT: Padigaru, Muralidhara
; APPLICANT: Rastelli, Luca
; APPLICANT: Spaderna, Steven
; APPLICANT: Shimkets, Richard
; APPLICANT: Zerhusen, Bryan
; APPLICANT: Spytek, Kimberly
; APPLICANT: Shenoy, Suresh
; APPLICANT: Li, Li
; APPLICANT: Gusev, Vladimir
; APPLICANT: Grosse, William
; APPLICANT: Alsobrook, John
; APPLICANT: Lepley, Denise
; APPLICANT: Burgess, Catherine
; APPLICANT: Gerlach, Valerie
; APPLICANT: Ellerman, Karen
; APPLICANT: MacDougall, John
; APPLICANT: Stone, David
; APPLICANT: Smithson, Glennda
; TITLE OF INVENTION: Novel Proteins and Nucleic Acids Encoding Same
; FILE REFERENCE: 21402-074 US
; CURRENT APPLICATION NUMBER: US/09/918,779
; CURRENT FILING DATE: 2001-07-30
; PRIOR APPLICATION NUMBER: 60/221,409
; PRIOR FILING DATE: 2000-07-28
; PRIOR APPLICATION NUMBER: 60/222,840
; PRIOR FILING DATE: 2000-08-04
; PRIOR APPLICATION NUMBER: 60/223,752
; PRIOR FILING DATE: 2000-08-08
; PRIOR APPLICATION NUMBER: 60/223,762
; PRIOR FILING DATE: 2000-08-08
; PRIOR APPLICATION NUMBER: 60/223,770
; PRIOR FILING DATE: 2000-08-08
; PRIOR APPLICATION NUMBER: 60/223,769
; PRIOR FILING DATE: 2000-08-08
; PRIOR APPLICATION NUMBER: 60/225,146
; PRIOR FILING DATE: 2000-08-14
; PRIOR APPLICATION NUMBER: 60/225,392
; PRIOR FILING DATE: 2000-08-15
; PRIOR APPLICATION NUMBER: 60/225,470
; PRIOR FILING DATE: 2000-08-15
; PRIOR APPLICATION NUMBER: 60/225,697
; PRIOR FILING DATE: 2000-08-16
; PRIOR APPLICATION NUMBER: 60/263,662
; PRIOR FILING DATE: 2001-02-01
; PRIOR APPLICATION NUMBER: 60/281,645
; PRIOR FILING DATE: 2001-04-05
; NUMBER OF SEQ ID NOS: 61
; SOFTWARE: PatentIn Ver. 2.1
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                  |||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
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                  |||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    2386 AACATCACCAAGGACACAAGGTTTGCTGAGCTGCTGGCTCTGGAGAGTGAAGCGGGGGTC 2445

Qy    2401 CCAGCCCTGGTGGGCCCCAGTGCCTTCAAGATCCCCTTCCTCATTCGGCAGAAGATAATT 2460
                  |||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    2446 CCAGCCCTGGTGGGCCCCAGTGCCTTCAAGATCCCCTTCCTCATTCGGCAGAAGATAATT 2505

Qy    2461 TCCAGCCTGGACCCACCCTGTAGGCGGGGTGCCGACTGGCGGACTCTGGCCCAGAAACTC 2520
                  |||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
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Db     2506 TCCAGCCTGGACCCACCCTGTAGGCGGGGTGCCGACTGGCGGACTCTGGCCCAGAAACTC 2565

Qy     2521 CACCTGGACAGCCATCTCAGCTTCTTTGCCTCCAAGCCCAGCCCCACAGCCATGATCCTC 2580
            ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db     2566 CACCTGGACAGCCATCTCAGCTTCTTTGCCTCCAAGCCCAGCCCCACAGCCATGATCCTC 2625

Qy     2581 AACCTGTGGGAGGCGCGGCACTTCCCCAACGGCAACCTCAGCCAGCTGGCTGCAGCAGTG 2640
            ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db     2626 AACCTGTGGGAGGCGCGGCACTTCCCCAACGGCAACCTCAGCCAGCTGGCTGCAGCAGTG 2685

Qy     2641 GCTGGACTGGGCCAGCCAGACGCTGGCCTCTTCACAGTGTCGGAGGCTGAGTGCTGA 2697
            ||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db     2686 GCTGGACTGGGCCAGCCAGACGCTGGCCTCTTCACAGTGTCGGAGGCTGAGTGCTGA 2742
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RESULT 2
US-10-624-932-1
; Sequence 1, Application US/10624932
; Publication No. US20040096877A1
; GENERAL INFORMATION:
; APPLICANT: Taupier, Raymond
; APPLICANT: Padigaru, Muralidhara
; APPLICANT: Rastelli, Luca
; APPLICANT: Spaderna, Steven
; APPLICANT: Shimkets, Richard
; APPLICANT: Zerhusen, Bryan
; APPLICANT: Spytek, Kimberly
; APPLICANT: Shenoy, Suresh
; APPLICANT: Li, Li
; APPLICANT: Gusev, Vladimir
; APPLICANT: Grosse, William
; APPLICANT: Alsobrook, John
; APPLICANT: Lepley, Denise
; APPLICANT: Burgess, Catherine
; APPLICANT: Gerlach, Valerie
; APPLICANT: Ellerman, Karen
; APPLICANT: MacDougall, John
; APPLICANT: Stone, David
; APPLICANT: Smithson, Glennda
; TITLE OF INVENTION: Novel Proteins and Nucleic Acids Encoding Same
; FILE REFERENCE: 21402-074 US
; CURRENT APPLICATION NUMBER: US/10/624,932
; CURRENT FILING DATE: 2003-07-21
; PRIOR APPLICATION NUMBER: 09/918,779
; PRIOR FILING DATE: 2001-07-03
; PRIOR APPLICATION NUMBER: 60/221,409
; PRIOR FILING DATE: 2000-07-28
; PRIOR APPLICATION NUMBER: 60/222,840
; PRIOR FILING DATE: 2000-08-04
; PRIOR APPLICATION NUMBER: 60/223,752
; PRIOR FILING DATE: 2000-08-08
; PRIOR APPLICATION NUMBER: 60/223,762
; PRIOR FILING DATE: 2000-08-08
; PRIOR APPLICATION NUMBER: 60/223,770
; PRIOR FILING DATE: 2000-08-08
; PRIOR APPLICATION NUMBER: 60/223,769
; PRIOR FILING DATE: 2000-08-08

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  Query Match              100.0%;   Score 2697;   DB 17;   Length 2752;
  Best Local Similarity    100.0%;   Pred. No. 0;
  Matches 2697;  Conservative    0;  Mismatches    0;  Indels    0;  Gaps    0;


Qy         1 ATGGCCGTCCGGCCCGGCCTGTGGCCAGCGCTCCTGGGCATAGTCCTCGCCGCTTGGCTC 60
             |||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        46 ATGGCCGTCCGGCCCGGCCTGTGGCCAGCGCTCCTGGGCATAGTCCTCGCCGCTTGGCTC 105


Qy        61 CGCGGCTCGGGTGCCCAGCAGAGTGCCACCGTGGCCAACCCAGTGCCTGGTGCCAACCCG 120
             |||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db       106 CGCGGCTCGGGTGCCCAGCAGAGTGCCACCGTGGCCAACCCAGTGCCTGGTGCCAACCCG 165


Qy       121 GACCTGCTTCCCCACTTCCTGGTGGAGCCCGAGGATGTGTACATCGTCAAGAACAAGCCA 180
             |||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db       166 GACCTGCTTCCCCACTTCCTGGTGGAGCCCGAGGATGTGTACATCGTCAAGAACAAGCCA 225


Qy       181 GTGCTGCTTGTGTGCAAGGCCGTGCCCGCCACGCAGATCTTCTTCAAGTGCAACGGGGAG 240
             |||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db       226 GTGCTGCTTGTGTGCAAGGCCGTGCCCGCCACGCAGATCTTCTTCAAGTGCAACGGGGAG 285


Qy       241 TGGGTGCGCCAGGTGGACCACGTGATCGAGCGCAGCACAGACGGGAGCAGTGGGCTGCCC 300
             |||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db       286 TGGGTGCGCCAGGTGGACCACGTGATCGAGCGCAGCACAGACGGGAGCAGTGGGCTGCCC 345


Qy       301 ACCATGGAGGTCCGCATTAATGTCTCAAGGCAGCAGGTCGAGAAGGTGTTCGGGCTGGAG 360
             |||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db       346 ACCATGGAGGTCCGCATTAATGTCTCAAGGCAGCAGGTCGAGAAGGTGTTCGGGCTGGAG 405


Qy       361 GAATACTGGTGCCAGTGCGTGGCATGGAGCTCCTCGGGCACCACCAAGAGTCAGAAGGCC 420
             |||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db       406 GAATACTGGTGCCAGTGCGTGGCATGGAGCTCCTCGGGCACCACCAAGAGTCAGAAGGCC 465


Qy       421 TACATCCGCATAGCCAGATTGCGCAAGAACTTCGAGCAGGAGCCGCTGGCCAAGGAGGTG 480
             |||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db       466 TACATCCGCATAGCCAGATTGCGCAAGAACTTCGAGCAGGAGCCGCTGGCCAAGGAGGTG 525


Qy       481 TCCCTGGAGCAGGGCATCGTGCTGCCCTGCCGTCCACCGGAGGGCATCCCTCCAGCCGAG 540
             |||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db       526 TCCCTGGAGCAGGGCATCGTGCTGCCCTGCCGTCCACCGGAGGGCATCCCTCCAGCCGAG 585


Qy       541 GTGGAGTGGCTCCGGAACGAGGACCTGGTGGACCCGTCCCTGGACCCCAATGTATACATC 600
             |||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
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Db    586  GTGGAGTGGCTCCGGAACGAGGACCTGGTGGACCCGTCCCTGGACCCCAATGTATACATC 645

Qy    601  ACGCGGGAGCACAGCCTGGTGGTGCGACAGGCCCGCCTTGCTGACACGGCCAACTACACC 660
           |||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    646  ACGCGGGAGCACAGCCTGGTGGTGCGACAGGCCCGCCTTGCTGACACGGCCAACTACACC 705

Qy    661  TGCGTGGCCAAGAACATCGTGGCACGTCGCCGCAGCGCCTCCGCTGCTGTCATCGTCTAC 720
           |||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    706  TGCGTGGCCAAGAACATCGTGGCACGTCGCCGCAGCGCCTCCGCTGCTGTCATCGTCTAC 765

Qy    721  GTGAACGGTGGGTGGTCGACGTGGACCGAGTGGTCCGTCTGCAGCGCCAGCTGTGGGCGC 780
           |||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    766  GTGAACGGTGGGTGGTCGACGTGGACCGAGTGGTCCGTCTGCAGCGCCAGCTGTGGGCGC 825

Qy    781  GGCTGGCAGAAACGGAGCCGGAGCTGCACCAACCCGGCGCCTCTCAACGGGGGCGCTTTC 840
           |||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    826  GGCTGGCAGAAACGGAGCCGGAGCTGCACCAACCCGGCGCCTCTCAACGGGGGCGCTTTC 885

Qy    841  TGTGAGGGGCAGAATGTCCAGAAAACAGCCTGCGCCACCCTGTGCCCAGTAGACGGCAGC 900
           |||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    886  TGTGAGGGGCAGAATGTCCAGAAAACAGCCTGCGCCACCCTGTGCCCAGTAGACGGCAGC 945

Qy    901  TGGAGCCCGTGGAGCAAGTGGTCGGCCTGTGGGCTGGACTGCACCCACTGGCGGAGCCGT 960
           |||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    946  TGGAGCCCGTGGAGCAAGTGGTCGGCCTGTGGGCTGGACTGCACCCACTGGCGGAGCCGT 1005

Qy    961  GAGTGCTCTGACCCAGCACCCCGCAACGGAGGGGAGGAGTGCCAGGGCACTGACCTGGAC 1020
           |||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db   1006  GAGTGCTCTGACCCAGCACCCCGCAACGGAGGGGAGGAGTGCCAGGGCACTGACCTGGAC 1065

Qy   1021  ACCCGCAACTGTACCAGTGACCTCTGTGTACACAGTGCTTCTGGCCCTGAGGACGTGGCC 1080
           |||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db   1066  ACCCGCAACTGTACCAGTGACCTCTGTGTACACAGTGCTTCTGGCCCTGAGGACGTGGCC 1125

Qy   1081  CTCTATGTGGGCCTCATCGCCGTGGCCGTCTGCCTGGTCCTGCTGCTGCTTGTCCTCATC 1140
           |||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db   1126  CTCTATGTGGGCCTCATCGCCGTGGCCGTCTGCCTGGTCCTGCTGCTGCTTGTCCTCATC 1185

Qy   1141  CTCGTTTATTGCCGGAAGAAGGAGGGGCTGGACTCAGATGTGGCTGACTCGTCCATTCTC 1200
           |||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db   1186  CTCGTTTATTGCCGGAAGAAGGAGGGGCTGGACTCAGATGTGGCTGACTCGTCCATTCTC 1245

Qy   1201  ACCTCAGGCTTCCAGCCCGTCAGCATCAAGCCCAGCAAAGCAGACAACCCCCATCTGCTC 1260
           |||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db   1246  ACCTCAGGCTTCCAGCCCGTCAGCATCAAGCCCAGCAAAGCAGACAACCCCCATCTGCTC 1305

Qy   1261  ACCATCCAGCCGGACCTCAGCACCACCACCACCACCTACCAGGGCAGTCTCTGTCCCCGG 1320
           |||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db   1306  ACCATCCAGCCGGACCTCAGCACCACCACCACCACCTACCAGGGCAGTCTCTGTCCCCGG 1365

Qy   1321  CAGGATGGGCCCAGCCCCAAGTTCCAGCTCACCAATGGGCACCTGCTCAGCCCCCTGGGT 1380
           |||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db   1366  CAGGATGGGCCCAGCCCCAAGTTCCAGCTCACCAATGGGCACCTGCTCAGCCCCCTGGGT 1425

Qy   1381  GGCGGCCGCCACACACTGCACCACAGCTCTCCCACCTCTGAGGCCGAGGAGTTCGTCTCC 1440
           |||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db   1426  GGCGGCCGCCACACACTGCACCACAGCTCTCCCACCTCTGAGGCCGAGGAGTTCGTCTCC 1485
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Qy   1441 CGCCTCTCCACCCAGAACTACTTCCGCTCCCTGCCCCGAGGCACCAGCAACATGACCTAT 1500
          |||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db   1486 CGCCTCTCCACCCAGAACTACTTCCGCTCCCTGCCCCGAGGCACCAGCAACATGACCTAT 1545

Qy   1501 GGGACCTTCAACTTCCTCGGGGGCCGGCTGATGATCCCTAATACAGGTATCAGCCTCCTC 1560
          |||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db   1546 GGGACCTTCAACTTCCTCGGGGGCCGGCTGATGATCCCTAATACAGGTATCAGCCTCCTC 1605

Qy   1561 ATCCCCCCAGATGCCATACCCCGAGGGAAGATCTATGAGATCTACCTCACGCTGCACAAG 1620
          |||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db   1606 ATCCCCCCAGATGCCATACCCCGAGGGAAGATCTATGAGATCTACCTCACGCTGCACAAG 1665

Qy   1621 CCGGAAGACGTGAGGTTGCCCCTAGCTGGCTGTCAGACCCTGCTGAGTCCCATCGTTAGC 1680
          |||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db   1666 CCGGAAGACGTGAGGTTGCCCCTAGCTGGCTGTCAGACCCTGCTGAGTCCCATCGTTAGC 1725

Qy   1681 TGTGGACCCCCTGGCGTCCTGCTCACCCGGCCAGTCATCCTGGCTATGGACCACTGTGGG 1740
          |||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db   1726 TGTGGACCCCCTGGCGTCCTGCTCACCCGGCCAGTCATCCTGGCTATGGACCACTGTGGG 1785

Qy   1741 GAGCCCAGCCCTGACAGCTGGAGCCTGCGCCTCAAAAAGCAGTCGTGCGAGGGCAGCTGG 1800
          |||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db   1786 GAGCCCAGCCCTGACAGCTGGAGCCTGCGCCTCAAAAAGCAGTCGTGCGAGGGCAGCTGG 1845

Qy   1801 GAGGATGTGCTGCACCTGGGCGAGGAGGCGCCCTCCCACCTCTACTACTGCCAGCTGGAG 1860
          |||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db   1846 GAGGATGTGCTGCACCTGGGCGAGGAGGCGCCCTCCCACCTCTACTACTGCCAGCTGGAG 1905

Qy   1861 GCCAGTGCCTGCTACGTCTTCACCGAGCAGCTGGGCCGCTTTGCCCTGGTGGGAGAGGCC 1920
          |||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db   1906 GCCAGTGCCTGCTACGTCTTCACCGAGCAGCTGGGCCGCTTTGCCCTGGTGGGAGAGGCC 1965

Qy   1921 CTCAGCGTGGCTGCCGCCAAGCGCCTCAAGCTGCTTCTGTTTGCGCCGGTGGCCTGCACC 1980
          |||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db   1966 CTCAGCGTGGCTGCCGCCAAGCGCCTCAAGCTGCTTCTGTTTGCGCCGGTGGCCTGCACC 2025

Qy   1981 TCCCTCGAGTACAACATCCGGGTCTACTGCCTGCATGACACCCACGATGCACTCAAGGAG 2040
          |||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db   2026 TCCCTCGAGTACAACATCCGGGTCTACTGCCTGCATGACACCCACGATGCACTCAAGGAG 2085

Qy   2041 GTGGTGCAGCTGGAGAAGCAGCTGGGGGGACAGCTGATCCAGGAGCCACGGGTCCTGCAC 2100
          |||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db   2086 GTGGTGCAGCTGGAGAAGCAGCTGGGGGGACAGCTGATCCAGGAGCCACGGGTCCTGCAC 2145

Qy   2101 TTCAAGGACAGTTACCACAACCTGCGCCTATCCATCCACGATGTGCCCAGCTCCCTGTGG 2160
          |||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db   2146 TTCAAGGACAGTTACCACAACCTGCGCCTATCCATCCACGATGTGCCCAGCTCCCTGTGG 2205

Qy   2161 AAGAGTAAGCTCCTTGTCAGCTACCAGGAGATCCCCTTTTATCACATCTGGAATGGCACG 2220
          |||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db   2206 AAGAGTAAGCTCCTTGTCAGCTACCAGGAGATCCCCTTTTATCACATCTGGAATGGCACG 2265

Qy   2221 CAGCGGTACTTGCACTGCACCTTCACCCTGGAGCGTGTCAGCCCCAGCACTAGTGACCTG 2280
          |||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db   2266 CAGCGGTACTTGCACTGCACCTTCACCCTGGAGCGTGTCAGCCCCAGCACTAGTGACCTG 2325
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Qy    2281 GCCTGCAAGCTGTGGGTGTGGCAGGTGGAGGGCGACGGGCAGAGCTTCAGCATCAACTTC 2340
           ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    2326 GCCTGCAAGCTGTGGGTGTGGCAGGTGGAGGGCGACGGGCAGAGCTTCAGCATCAACTTC 2385

Qy    2341 AACATCACCAAGGACACAAGGTTTGCTGAGCTGCTGGCTCTGGAGAGTGAAGCGGGGGTC 2400
           ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    2386 AACATCACCAAGGACACAAGGTTTGCTGAGCTGCTGGCTCTGGAGAGTGAAGCGGGGGTC 2445

Qy    2401 CCAGCCCTGGTGGGCCCCAGTGCCTTCAAGATCCCCTTCCTCATTCGGCAGAAGATAATT 2460
           ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    2446 CCAGCCCTGGTGGGCCCCAGTGCCTTCAAGATCCCCTTCCTCATTCGGCAGAAGATAATT 2505

Qy    2461 TCCAGCCTGGACCCACCCTGTAGGCGGGGTGCCGACTGGCGGACTCTGGCCCAGAAACTC 2520
           ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    2506 TCCAGCCTGGACCCACCCTGTAGGCGGGGTGCCGACTGGCGGACTCTGGCCCAGAAACTC 2565

Qy    2521 CACCTGGACAGCCATCTCAGCTTCTTTGCCTCCAAGCCCAGCCCCACAGCCATGATCCTC 2580
           ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    2566 CACCTGGACAGCCATCTCAGCTTCTTTGCCTCCAAGCCCAGCCCCACAGCCATGATCCTC 2625

Qy    2581 AACCTGTGGGAGGCGCGGCACTTCCCCAACGGCAACCTCAGCCAGCTGGCTGCAGCAGTG 2640
           ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    2626 AACCTGTGGGAGGCGCGGCACTTCCCCAACGGCAACCTCAGCCAGCTGGCTGCAGCAGTG 2685

Qy    2641 GCTGGACTGGGCCAGCCAGACGCTGGCCTCTTCACAGTGTCGGAGGCTGAGTGCTGA 2697
           |||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    2686 GCTGGACTGGGCCAGCCAGACGCTGGCCTCTTCACAGTGTCGGAGGCTGAGTGCTGA 2742
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RESULT 3
US-09-970-944-1
; Sequence 1, Application US/09970944
; Publication No. US20030204052A1
; GENERAL INFORMATION:
; APPLICANT: Herrman, John L
; APPLICANT: Rastelli, Luca
; APPLICANT: Shimkets, Richard A
; TITLE OF INVENTION: No. US20030204052A1el Proteins and Nucleic Acids Encoding
Same and
; TITLE OF INVENTION: Antibodies Directed Against these Proteins
; FILE REFERENCE: 21402-138
; CURRENT APPLICATION NUMBER: US/09/970,944
; CURRENT FILING DATE: 2002-05-02
; PRIOR APPLICATION NUMBER: 60/237,862
; PRIOR FILING DATE: 2000-10-04
; NUMBER OF SEQ ID NOS: 62
; SOFTWARE: PatentIn Ver. 2.1
; SEQ ID NO 1
;   LENGTH: 2881
;   TYPE: DNA
;   ORGANISM: Homo sapiens
US-09-970-944-1

  Query Match   97.2%; Score 2621.4; DB 10; Length 2881;
  Best Local Similarity   98.9%; Pred. No. 0;
  Matches 2673; Conservative   0; Mismatches   21; Indels   9; Gaps   3;

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Qy      1 ATGGCCGTCCGGCCCGGCCTGTGGCCAGCGCTCCTGGGCATAGTCCTCGCCGCTTGGCTC 60
          |||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db     87 ATGGCCGTCCGGCCCGGCCTGTGGCCAGCGCTCCTGGGCATAGTCCTCGCCGCTTGGCTC 146

Qy     61 CGCGGCTCGGGTGCCCAGCAGAGTGCCACCGTGGCCAACCCAGTGCCTGGTGCCAACCCG 120
          |||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    147 CGCGGCTCGGGTGCCCAGCAGAGTGCCACCGTGGCCAACCCAGTGCCTGGTGCCAACCCG 206

Qy    121 GACCTGCTTCCCCACTTCCTGGTGGAGCCCGAGGATGTGTACATCGTCAAGAACAAGCCA 180
          |||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    207 GACCTGCTTCCCCACTTCCTGGTGGAGCCCGAGGATGTGTACATCGTCAAGAACAAGCCA 266

Qy    181 GTGCTGCTTGTGTGCAAGGCCGTGCCCGCCACGCAGATCTTCTTCAAGTGCAACGGGGAG 240
          |||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    267 GTGCTGCTTGTGTGCAAGGCCGTGCCCGCCACGCAGATCTTCTTCAAGTGCAACGGGGAG 326

Qy    241 TGGGTGCGCCAGGTGGACCACGTGATCGAGCGCAGCACAGACGGGAGCAGTGGGCTGCCC 300
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||   |||
Db    327 TGGGTGCGCCAGGTGGACCACGTGATCGAGCGCAGCACAGACGGGAGCAGTGGTGAGCCG 386

Qy    301 ACCATGGAGGTCCGCATTAATGTCTCAAGGCAGCAGGTCGAGAAGGTGTTCGGGCTGGAG 360
          |||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    387 ACCATGGAGGTCCGCATTAATGTCTCAAGGCAGCAGGTCGAGAAGGTGTTCGGGCTGGAG 446

Qy    361 GAATACTGGTGCCAGTGCGTGGCATGGAGCTCCTCGGGCACCACCAAGAGTCAGAAGGCC 420
          |||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    447 GAATACTGGTGCCAGTGCGTGGCATGGAGCTCCTCGGGCACCACCAAGAGTCAGAAGGCC 506

Qy    421 TACATCCGCATAGCCAGATTGCGCAAGAACTTCGAGCAGGAGCCGCTGGCCAAGGAGGTG 480
          |||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    507 TACATCCGCATAGCCAGATTGCGCAAGAACTTCGAGCAGGAGCCGCTGGCCAAGGAGGTG 566

Qy    481 TCCCTGGAGCAGGGCATCGTGCTGCCCTGCCGTCCACCGGAGGGCATCCCTCCAGCCGAG 540
          |||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    567 TCCCTGGAGCAGGGCATCGTGCTGCCCTGCCGTCCACCGGAGGGCATCCCTCCAGCCGAG 626

Qy    541 GTGGAGTGGCTCCGGAACGAGGACCTGGTGGACCCGTCCCTGGACCCCAATGTATACATC 600
          |||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    627 GTGGAGTGGCTCCGGAACGAGGACCTGGTGGACCCGTCCCTGGACCCCAATGTATACATC 686

Qy    601 ACGCGGGAGCACAGCCTGGTGGTGCGACAGGCCCGCCTTGCTGACACGGCCAACTACACC 660
          |||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    687 ACGCGGGAGCACAGCCTGGTGGTGCGACAGGCCCGCCTTGCTGACACGGCCAACTACACC 746

Qy    661 TGCGTGGCCAAGAACATCGTGGCACGTCGCCGCAGCGCCTCCGCTGCTGTCATCGTCTAC 720
          ||||||||||||||||||||||||||.-|||||||||||||||||||||||||||||||||
Db    747 TGCGTGGCCAAGAACATCGTGGCACGTCGCCGCAGCGCCTCCGCTGCTGTCATCGTCTAC 806

Qy    721 GTGAACGGTGGGTGGTCGACGTGGACCGAGTGGTCCGTCTGCAGCGCCAGCTGTGGGCGC 780
          |||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    807 GTGAACGGTGGGTGGTCGACGTGGACCGAGTGGTCCGTCTGCAGCGCCAGCTGTGGGCGC 866

Qy    781 GGCTGGCAGAAACGGAGCCGGAGCTGCACCAACCCGGCGCCTCTCAACGGGGGCGCTTTC 840
          |||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    867 GGCTGGCAGAAACGGAGCCGGAGCTGCACCAACCCGGCGCCTCTCAACGGGGGCGCTTTC 926

Qy    841 TGTGAGGGGCAGAATGTCCAGAA---AACAGCCTGCGCCACCCTGTGCCCAGTAGACGGC 897
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            |||||||||||||||||||||||  |     ||  |  ||  |  |      ||    |  |  ||  ||||||
Db    927 TGTGAGGGGCAGAATGTCCATGACCGCACCGTCTCCTCTCTGCTTGTCTCTGTGGACGGC 986

Qy    898 AGCTGGAGCCCGTGGAGCAAGTGGTCGGCCTGTGGGCTGGACTGCACCCACTGGCGGAGC 957
            ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    987 AGCTGGAGCCCGTGGAGCAAGTGGTCGGCCTGTGGGCTGGACTGCACCCACTGGCGGAGC 1046

Qy    958 CGTGAGTGCTCTGACCCAGCACCCCGCAACGGAGGGGAGGAGTGCCAGGGCACTGACCTG 1017
            ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db   1047 CGTGAGTGCTCTGACCCAGCACCCCGCAACGGAGGGGAGGAGTGCCAGGGCACTGACCTG 1106

Qy   1018 GACACCCGCAACTGTACCAGTGACCTCTGTGTACACAGTGCTTCTGGCCCTGAGGACGTG 1077
            ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db   1107 GACACCCGCAACTGTACCAGTGACCTCTGTGTACACAGTGCTTCTGGCCCTGAGGACGTG 1166

Qy   1078 GCCCTCTATGTGGGCCTCATCGCCGTGGCCGTCTGCCTGGTCCTGCTGCTGCTTGTCCTC 1137
            ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db   1167 GCCCTCTATGTGGGCCTCATCGCCGTGGCCGTCTGCCTGGTCCTGCTGCTGCTTGTCCTC 1226

Qy   1138 ATCCTCGTTTATTGCCGGAAGAAGGAGGGGCTGGACTCAGATGTGGCTGACTCGTCCATT 1197
            ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db   1227 ATCCTCGTTTATTGCCGGAAGAAGGAGGGGCTGGACTCAGATGTGGCTGACTCGTCCATT 1286

Qy   1198 CTCACCTCAGGCTTCCAGCCCGTCAGCATCAAGCCCAGCAAAGCAGACAACCCCCATCTG 1257
            ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db   1287 CTCACCTCAGGCTTCCAGCCCGTCAGCATCAAGCCCAGCAAAGCAGACAACCCCCATCTG 1346

Qy   1258 CTCACCATCCAGCCGGACCTCAGCACCACCACCACCACCTACCAGGGCAGTCTCTGTCCC 1317
            |||||||||||||||||||||       |||||||||||||||||||||||||||||||||
Db   1347 CTCACCATCCAGCCGGACCTCAG---CACCACCACCACCTACCAGGGCAGTCTCTGTCCC 1403

Qy   1318 CGGCAGGATGGGCCCAGCCCCAAGTTCCAGCTCACCAATGGGCACCTGCTCAGCCCCCTG 1377
            ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db   1404 CGGCAGGATGGGCCCAGCCCCAAGTTCCAGCTCACCAATGGGCACCTGCTCAGCCCCCTG 1463

Qy   1378 GGTGGCGGCCGCCACACACTGCACCACAGCTCTCCCACCTCTGAGGCCGAGGAGTTCGTC 1437
            ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db   1464 GGTGGCGGCCGCCACACACTGCACCACAGCTCTCCCACCTCTGAGGCCGAGGAGTTCGTC 1523

Qy   1438 TCCCGCCTCTCCACCCAGAACTACTTCCGCTCCCTGCCCCGAGGCACCAGCAACATGACC 1497
            ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db   1524 TCCCGCCTCTCCACCCAGAACTACTTCCGCTCCCTGCCCCGAGGCACCAGCAACATGACC 1583

Qy   1498 TATGGGACCTTCAACTTCCTCGGGGGCCGGCTGATGATCCCTAATACAGGTATCAGCCTC 1557
            ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db   1584 TATGGGACCTTCAACTTCCTCGGGGGCCGGCTGATGATCCCTAATACAGGTATCAGCCTC 1643

Qy   1558 CTCATCCCCCCAGATGCCATACCCCGAGGGAAGATCTATGAGATCTACCTCACGCTGCAC 1617
            ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db   1644 CTCATCCCCCCAGATGCCATACCCCGAGGGAAGATCTATGAGATCTACCTCACGCTGCAC 1703

Qy   1618 AAGCCGGAAGACGTGAGGTTGCCCCTAGCTGGCTGTCAGACCCTGCTGAGTCCCATCGTT 1677
            ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db   1704 AAGCCGGAAGACGTGAGGTTGCCCCTAGCTGGCTGTCAGACCCTGCTGAGTCCCATCGTT 1763

Qy   1678 AGCTGTGGACCCCCTGGCGTCCTGCTCACCCGGCCAGTCATCCTGGCTATGGACCACTGT 1737
            ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
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Db    1764 AGCTGTGGACCCCCTGGCGTCCTGCTCACCCGGCCAGTCATCCTGGCTATGGACCACTGT 1823

Qy    1738 GGGGAGCCCAGCCCTGACAGCTGGAGCCTGCGCCTCAAAAAGCAGTCGTGCGAGGGCAGC 1797
           ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    1824 GGGGAGCCCAGCCCTGACAGCTGGAGCCTGCGCCTCAAAAAGCAGTCGTGCGAGGGCAGC 1883

Qy    1798 TGGG---AGGATGTGCTGCACCTGGGCGAGGAGGCGCCCTCCCACCTCTACTACTGCCAG 1854
           ||||   |||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    1884 TGGGAGCAGGATGTGCTGCACCTGGGCGAGGAGGCGCCCTCCCACCTCTACTACTGCCAG 1943

Qy    1855 CTGGAGGCCAGTGCCTGCTACGTCTTCACCGAGCAGCTGGGCCGCTTTGCCCTGGTGGGA 1914
           ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    1944 CTGGAGGCCAGTGCCTGCTACGTCTTCACCGAGCAGCTGGGCCGCTTTGCCCTGGTGGGA 2003

Qy    1915 GAGGCCCTCAGCGTGGCTGCCGCCAAGCGCCTCAAGCTGCTTCTGTTTGCGCCGGTGGCC 1974
           ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    2004 GAGGCCCTCAGCGTGGCTGCCGCCAAGCGCCTCAAGCTGCTTCTGTTTGCGCCGGTGGCC 2063

Qy    1975 TGCACCTCCCTCGAGTACAACATCCGGGTCTACTGCCTGCATGACACCCACGATGCACTC 2034
           ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    2064 TGCACCTCCCTCGAGTACAACATCCGGGTCTACTGCCTGCATGACACCCACGATGCACTC 2123

Qy    2035 AAGGAGGTGGTGCAGCTGGAGAAGCAGCTGGGGGGACAGCTGATCCAGGAGCCACGGGTC 2094
           ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    2124 AAGGAGGTGGTGCAGCTGGAGAAGCAGCTGGGGGGACAGCTGATCCAGGAGCCACGGGTC 2183

Qy    2095 CTGCACTTCAAGGACAGTTACCACAACCTGCGCCTATCCATCCACGATGTGCCCAGCTCC 2154
           ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    2184 CTGCACTTCAAGGACAGTTACCACAACCTGCGCCTATCCATCCACGATGTGCCCAGCTCC 2243

Qy    2155 CTGTGGAAGAGTAAGCTCCTTGTCAGCTACCAGGAGATCCCCTTTTATCACATCTGGAAT 2214
           ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    2244 CTGTGGAAGAGTAAGCTCCTTGTCAGCTACCAGGAGATCCCCTTTTATCACATCTGGAAT 2303

Qy    2215 GGCACGCAGCGGTACTTGCACTGCACCTTCACCCTGGAGCGTGTCAGCCCCAGCACTAGT 2274
           |||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    2304 GGCACGCAGCGGTACTTGCACTGCACCTTCACCCTGGAGCGTGTCAGCCCCAGCACTAGT 2363

Qy    2275 GACCTGGCCTGCAAGCTGTGGGTGTGGCAGGTGGAGGGCGACGGGCAGAGCTTCAGCATC 2334
           ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    2364 GACCTGGCCTGCAAGCTGTGGGTGTGGCAGGTGGAGGGCGACGGGCAGAGCTTCAGCATC 2423

Qy    2335 AACTTCAACATCACCAAGGACACAAGGTTTGCTGAGCTGCTGGCTCTGGAGAGTGAAGCG 2394
           ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    2424 AACTTCAACATCACCAAGGACACAAGGTTTGCTGAGCTGCTGGCTCTGGAGAGTGAAGCG 2483

Qy    2395 GGGGTCCCAGCCCTGGTGGGCCCCAGTGCCTTCAAGATCCCCTTCCTCATTCGGCAGAAG 2454
           ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    2484 GGGGTCCCAGCCCTGGTGGGCCCCAGTGCCTTCAAGATCCCCTTCCTCATTCGGCAGAAG 2543

Qy    2455 ATAATTTCCAGCCTGGACCCACCCTGTAGGCGGGGTGCCGACTGGCGGACTCTGGCCCAG 2514
           ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    2544 ATAATTTCCAGCCTGGACCCACCCTGTAGGCGGGGTGCCGACTGGCGGACTCTGGCCCAG 2603

Qy    2515 AAACTCCACCTGGACAGCCATCTCAGCTTCTTTGCCTCCAAGCCCAGCCCCACAGCCATG 2574
           ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    2604 AAACTCCACCTGGACAGCCATCTCAGCTTCTTTGCCTCCAAGCCCAGCCCCACAGCCATG 2663
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Qy     2575 ATCCTCAACCTGTGGGAGGCGCGGCACTTCCCCAACGGCAACCTCAGCCAGCTGGCTGCA 2634
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Db     2664 ATCCTCAACCTGTGGGAGGCGCGGCACTTCCCCAACGGCAACCTCAGCCAGCTGGCTGCA 2723

Qy     2635 GCAGTGGCTGGACTGGGCCAGCCAGACGCTGGCCTCTTCACAGTGTCGGAGGCTGAGTGC 2694
            ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
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Qy     2695 TGA 2697
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Db     2784 TGA 2786
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RESULT 4
US-10-643-795A-77
; Sequence 77, Application US/10643795A
; Publication No. US20040241703A1
; GENERAL INFORMATION:
; APPLICANT: FREDERIC J. DESAUVAGE
; APPLICANT: GRETCHEN FRANTZ
; APPLICANT: KENNETH J. HILLAN
; APPLICANT: PAUL POLAKIS
; APPLICANT: ANDREW POLSON
; APPLICANT: VICTORIA SMITH
; APPLICANT: SUSAN D. SPENCER
; APPLICANT: THOMAS D. WU
; APPLICANT: ZEMIN ZHANG
; TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR THE DIAGNOSIS AND
; TITLE OF INVENTION: TREATMENT OF TUMOR
; FILE REFERENCE: P5026R1-US
; CURRENT APPLICATION NUMBER: US/10/643,795A
; CURRENT FILING DATE: 2003-08-19
; PRIOR APPLICATION NUMBER: US 60/404,809
; PRIOR FILING DATE: 2002-08-19
; PRIOR APPLICATION NUMBER: US 60/405,645
; PRIOR FILING DATE: 2002-08-21
; PRIOR APPLICATION NUMBER: US 60/413,192
; PRIOR FILING DATE: 2002-09-23
; PRIOR APPLICATION NUMBER: US 60/419,008
; PRIOR FILING DATE: 2002-10-15
; PRIOR APPLICATION NUMBER: US 60/426,847
; PRIOR FILING DATE: 2002-11-15
; PRIOR APPLICATION NUMBER: US 60/484,959
; PRIOR FILING DATE: 2003-07-02
; NUMBER OF SEQ ID NOS: 158
; SEQ ID NO 77
;   LENGTH: 3561
;   TYPE: DNA
;   ORGANISM: Homo sapien
US-10-643-795A-77

  Query Match   86.9%; Score 2343; DB 18; Length 3561;
  Best Local Similarity 93.6%; Pred. No. 0;
  Matches 2524; Conservative   0; Mismatches   5; Indels 168; Gaps   1;

Qy   1 ATGGCCGTCCGGCCCGGCCTGTGGCCAGCGCTCCTGGGCATAGTCCTCGCCGCTTGGCTC 60
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Db         4 ATGGCCGTCCGGCCCGGCCTGTGGCCAGCGCTCCTGGGCATAGTCCTCGCCGCTTGGCTC 63

Qy        61 CGCGGCTCGGGTGCCCAGCAGAGTGCCACCGTGGCCAACCCAGTGCCTGGTGCCAACCCG 120
                    ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        64 CGCGGCTCGGGTGCCCAGCAGAGTGCCACCGTGGCCAACCCAGTGCCTGGTGCCAACCCG 123

Qy       121 GACCTGCTTCCCCACTTCCTGGTGGAGCCCGAGGATGTGTACATCGTCAAGAACAAGCCA 180
                    ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db       124 GACCTGCTTCCCCACTTCCTGGTGGAGCCCGAGGATGTGTACATCGTCAAGAACAAGCCA 183

Qy       181 GTGCTGCTTGTGTGCAAGGCCGTGCCCGCCACGCAGATCTTCTTCAAGTGCAACGGGGAG 240
                    ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db       184 GTGCTGCTTGTGTGCAAGGCCGTGCCCGCCACGCAGATCTTCTTCAAGTGCAACGGGGAG 243

Qy       241 TGGGTGCGCCAGGTGGACCACGTGATCGAGCGCAGCACAGACGGGAGCAGTGGGCTGCCC 300
                    ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db       244 TGGGTGCGCCAGGTGGACCACGTGATCGAGCGCAGCACAGACGGGAGCAGTGGGCTGCCC 303

Qy       301 ACCATGGAGGTCCGCATTAATGTCTCAAGGCAGCAGGTCGAGAAGGTGTTCGGGCTGGAG 360
                    ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db       304 ACCATGGAGGTCCGCATTAATGTCTCAAGGCAGCAGGTCGAGAAGGTGTTCGGGCTGGAG 363

Qy       361 GAATACTGGTGCCAGTGCGTGGCATGGAGCTCCTCGGGCACCACCAAGAGTCAGAAGGCC 420
                    ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db       364 GAATACTGGTGCCAGTGCGTGGCATGGAGCTCCTCGGGCACCACCAAGAGTCAGAAGGCC 423

Qy       421 TACATCCGCATAGCCAGATTGCGCAAGAACTTCGAGCAGGAGCCGCTGGCCAAGGAGGTG 480
                    ||||||||||||||||   |||||||||||||||||||||||||||||||||||||||||
Db       424 TACATCCGCATAGCCTATTTGCGCAAGAACTTCGAGCAGGAGCCGCTGGCCAAGGAGGTG 483

Qy       481 TCCCTGGAGCAGGGCATCGTGCTGCCCTGCCGTCCACCGGAGGGCATCCCTCCAGCCGAG 540
                    ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db       484 TCCCTGGAGCAGGGCATCGTGCTGCCCTGCCGTCCACCGGAGGGCATCCCTCCAGCCGAG 543

Qy       541 GTGGAGTGGCTCCGGAACGAGGACCTGGTGGACCCGTCCCTGGACCCCAATGTATACATC 600
                    ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db       544 GTGGAGTGGCTCCGGAACGAGGACCTGGTGGACCCGTCCCTGGACCCCAATGTATACATC 603

Qy       601 ACGCGGGAGCACAGCCTGGTGGTGCGACAGGCCCGCCTTGCTGACACGGCCAACTACACC 660
                    ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db       604 ACGCGGGAGCACAGCCTGGTGGTGCGACAGGCCCGCCTTGCTGACACGGCCAACTACACC 663

Qy       661 TGCGTGGCCAAGAACATCGTGGCACGTCGCCGCAGCGCCTCCGCTGCTGTCATCGTCTAC 720
                    ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db       664 TGCGTGGCCAAGAACATCGTGGCACGTCGCCGCAGCGCCTCCGCTGCTGTCATCGTCTAC 723

Qy       721 GTGAACGGTGGGTGGTCGACGTGGACCGAGTGGTCCGTCTGCAGCGCCAGCTGTGGGCGC 780
                    |||
Db       724 GTG--------------------------------------------------------- 726

Qy       781 GGCTGGCAGAAACGGAGCCGGAGCTGCACCAACCCGGCGCCTCTCAACGGGGGCGCTTTC 840

Db       727 --------------------------------------------------------- 726

Qy       841 TGTGAGGGGCAGAATGTCCAGAAAACAGCCTGCGCCACCCTGTGCCCAGTAGACGGCAGC 900
                                                                  |||||||||
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Db      727 ---------------------------------------------------GACGGCAGC 735

Qy      901 TGGAGCCCGTGGAGCAAGTGGTCGGCCTGTGGGCTGGACTGCACCCACTGGCGGAGCCGT 960
            |||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db      736 TGGAGCCCGTGGAGCAAGTGGTCGGCCTGTGGGCTGGACTGCACCCACTGGCGGAGCCGT 795

Qy      961 GAGTGCTCTGACCCAGCACCCCGCAACGGAGGGGAGGAGTGCCAGGGCACTGACCTGGAC 1020
            |||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db      796 GAGTGCTCTGACCCAGCACCCCGCAACGGAGGGGAGGAGTGCCAGGGCACTGACCTGGAC 855

Qy     1021 ACCCGCAACTGTACCAGTGACCTCTGTGTACACAGTGCTTCTGGCCCTGAGGACGTGGCC 1080
            ||||||||||||||||||||||||||||||||||| |||||||||||||||||||||||
Db      856 ACCCGCAACTGTACCAGTGACCTCTGTGTACACACTGCTTCTGGCCCTGAGGACGTGGCC 915

Qy     1081 CTCTATGTGGGCCTCATCGCCGTGGCCGTCTGCCTGGTCCTGCTGCTGCTTGTCCTCATC 1140
            |||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db      916 CTCTATGTGGGCCTCATCGCCGTGGCCGTCTGCCTGGTCCTGCTGCTGCTTGTCCTCATC 975

Qy     1141 CTCGTTTATTGCCGGAAGAAGGAGGGGCTGGACTCAGATGTGGCTGACTCGTCCATTCTC 1200
            |||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db      976 CTCGTTTATTGCCGGAAGAAGGAGGGGCTGGACTCAGATGTGGCTGACTCGTCCATTCTC 1035

Qy     1201 ACCTCAGGCTTCCAGCCCGTCAGCATCAAGCCCAGCAAAGCAGACAACCCCCATCTGCTC 1260
            |||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db     1036 ACCTCAGGCTTCCAGCCCGTCAGCATCAAGCCCAGCAAAGCAGACAACCCCCATCTGCTC 1095

Qy     1261 ACCATCCAGCCGGACCTCAGCACCACCACCACCACCTACCAGGGCAGTCTCTGTCCCCGG 1320
            |||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db     1096 ACCATCCAGCCGGACCTCAGCACCACCACCACCACCTACCAGGGCAGTCTCTGTCCCCGG 1155

Qy     1321 CAGGATGGGCCCAGCCCCAAGTTCCAGCTCACCAATGGGCACCTGCTCAGCCCCCTGGGT 1380
            |||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db     1156 CAGGATGGGCCCAGCCCCAAGTTCCAGCTCACCAATGGGCACCTGCTCAGCCCCCTGGGT 1215

Qy     1381 GGCGGCCGCCACACACTGCACCACAGCTCTCCCACCTCTGAGGCCGAGGAGTTCGTCTCC 1440
            |||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db     1216 GGCGGCCGCCACACACTGCACCACAGCTCTCCCACCTCTGAGGCCGAGGAGTTCGTCTCC 1275

Qy     1441 CGCCTCTCCACCCAGAACTACTTCCGCTCCCTGCCCCGAGGCACCAGCAACATGACCTAT 1500
            |||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db     1276 CGCCTCTCCACCCAGAACTACTTCCGCTCCCTGCCCCGAGGCACCAGCAACATGACCTAT 1335

Qy     1501 GGGACCTTCAACTTCCTCGGGGGCCGGCTGATGATCCCTAATACAGGTATCAGCCTCCTC 1560
            ||||||||||||||||||||||||||||||||||||||||||| |||||||||||
Db     1336 GGGACCTTCAACTTCCTCGGGGGCCGGCTGATGATCCCTAATACAGGAATCAGCCTCCTC 1395

Qy     1561 ATCCCCCCAGATGCCATACCCCGAGGGAAGATCTATGAGATCTACCTCACGCTGCACAAG 1620
            |||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db     1396 ATCCCCCCAGATGCCATACCCCGAGGGAAGATCTATGAGATCTACCTCACGCTGCACAAG 1455

Qy     1621 CCGGAAGACGTGAGGTTGCCCCTAGCTGGCTGTCAGACCCTGCTGAGTCCCATCGTTAGC 1680
            |||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db     1456 CCGGAAGACGTGAGGTTGCCCCTAGCTGGCTGTCAGACCCTGCTGAGTCCCATCGTTAGC 1515

Qy     1681 TGTGGACCCCCTGGCGTCCTGCTCACCCGGCCAGTCATCCTGGCTATGGACCACTGTGGG 1740
            ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db     1516 TGTGGACCCCCTGGCGTCCTGCTCACCCGGCCAGTCATCCTGGCTATGGACCACTGTGGG 1575
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Qy     1741 GAGCCCAGCCCTGACAGCTGGAGCCTGCGCCTCAAAAAGCAGTCGTGCGAGGGCAGCTGG 1800
            ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db     1576 GAGCCCAGCCCTGACAGCTGGAGCCTGCGCCTCAAAAAGCAGTCGTGCGAGGGCAGCTGG 1635

Qy     1801 GAGGATGTGCTGCACCTGGGCGAGGAGGCGCCCTCCCACCTCTACTACTGCCAGCTGGAG 1860
            ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db     1636 GAGGATGTGCTGCACCTGGGCGAGGAGGCGCCCTCCCACCTCTACTACTGCCAGCTGGAG 1695

Qy     1861 GCCAGTGCCTGCTACGTCTTCACCGAGCAGCTGGGCCGCTTTGCCCTGGTGGGAGAGGCC 1920
            ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db     1696 GCCAGTGCCTGCTACGTCTTCACCGAGCAGCTGGGCCGCTTTGCCCTGGTGGGAGAGGCC 1755

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            ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db     1756 CTCAGCGTGGCTGCCGCCAAGCGCCTCAAGCTGCTTCTGTTTGCGCCGGTGGCCTGCACC 1815

Qy     1981 TCCCTCGAGTACAACATCCGGGTCTACTGCCTGCATGACACCCACGATGCACTCAAGGAG 2040
            ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db     1816 TCCCTCGAGTACAACATCCGGGTCTACTGCCTGCATGACACCCACGATGCACTCAAGGAG 1875

Qy     2041 GTGGTGCAGCTGGAGAAGCAGCTGGGGGGACAGCTGATCCAGGAGCCACGGGTCCTGCAC 2100
            ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    .1876 GTGGTGCAGCTGGAGAAGCAGCTGGGGGGACAGCTGATCCAGGAGCCACGGGTCCTGCAC 1935

Qy     2101 TTCAAGGACAGTTACCACAACCTGCGCCTATCCATCCACGATGTGCCCAGCTCCCTGTGG 2160
            ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db     1936 TTCAAGGACAGTTACCACAACCTGCGCCTATCCATCCACGATGTGCCCAGCTCCCTGTGG 1995

Qy     2161 AAGAGTAAGCTCCTTGTCAGCTACCAGGAGATCCCCTTTTATCACATCTGGAATGGCACG 2220
            ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db     1996 AAGAGTAAGCTCCTTGTCAGCTACCAGGAGATCCCCTTTTATCACATCTGGAATGGCACG 2055

Qy     2221 CAGCGGTACTTGCACTGCACCTTCACCCTGGAGCGTGTCAGCCCCAGCACTAGTGACCTG 2280
            ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db     2056 CAGCGGTACTTGCACTGCACCTTCACCCTGGAGCGTGTCAGCCCCAGCACTAGTGACCTG 2115

Qy     2281 GCCTGCAAGCTGTGGGTGTGGCAGGTGGAGGGCGACGGGCAGAGCTTCAGCATCAACTTC 2340
            ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db     2116 GCCTGCAAGCTGTGGGTGTGGCAGGTGGAGGGCGACGGGCAGAGCTTCAGCATCAACTTC 2175

Qy     2341 AACATCACCAAGGACACAAGGTTTGCTGAGCTGCTGGCTCTGGAGAGTGAAGCGGGGGTC 2400
            ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db     2176 AACATCACCAAGGACACAAGGTTTGCTGAGCTGCTGGCTCTGGAGAGTGAAGCGGGGGTC 2235

Qy     2401 CCAGCCCTGGTGGGCCCCAGTGCCTTCAAGATCCCCTTCCTCATTCGGCAGAAGATAATT 2460
            ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db     2236 CCAGCCCTGGTGGGCCCCAGTGCCTTCAAGATCCCCTTCCTCATTCGGCAGAAGATAATT 2295

Qy     2461 TCCAGCCTGGACCCACCCTGTAGGCGGGGTGCCGACTGGCGGACTCTGGCCCAGAAACTC 2520
            ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db     2296 TCCAGCCTGGACCCACCCTGTAGGCGGGGTGCCGACTGGCGGACTCTGGCCCAGAAACTC 2355

Qy     2521 CACCTGGACAGCCATCTCAGCTTCTTTGCCTCCAAGCCCAGCCCCACAGCCATGATCCTC 2580
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Db     2356 CACCTGGACAGCCATCTCAGCTTCTTTGCCTCCAAGCCCAGCCCCACAGCCATGATCCTC 2415
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Qy   2581 AACCTGTGGGAGGCGCGGCACTTCCCCAACGGCAACCTCAGCCAGCTGGCTGCAGCAGTG 2640
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Db   2416 AACCTGTGGGAGGCGCGGCACTTCCCCAACGGCAACCTCAGCCAGCTGGCTGCAGCAGTG 2475

Qy   2641 GCTGGACTGGGCCAGCCAGACGCTGGCCTCTTCACAGTGTCGGAGGCTGAGTGCTGA 2697
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Db   2476 GCTGGACTGGGCCAGCCAGACGCTGGCCTCTTCACAGTGTCGGAGGCTGAGTGCTGA 2532
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RESULT 5
US-10-311-623-13
; Sequence 13, Application US/10311623
; Publication No. US20040023244A1
; GENERAL INFORMATION:
; APPLICANT: INCYTE GENOMICS, INC.; GRIFFIN, Jennifer A.
; APPLICANT: KALLICK, Deborah A.; TRIBOULEY, Catherine M.
; APPLICANT: YUE, Henry; NGUYEN, Danniel B.
; APPLICANT: TANG, Y. Tom; LAL, Preeti G.
; APPLICANT: POLICKY, Jennifer L.; AZIMZAI, Yalda
; APPLICANT: LU, Dyung Aina M.; GRAUL, Richard C.
; APPLICANT: YAO, Monique G.; BURFORD, Neil
; APPLICANT: HAFALIA, April J. A.; BAUGHN, Mariah R.
; APPLICANT: BANDMAN, Olga; ARVIZU, Chandra S.
; APPLICANT: YANG, Junming; XU, Yuming
; APPLICANT: GANDHI, Ameena R.; WARREN, Bridget A.
; APPLICANT: DING, Li; SANJANWALA, Madhusudan M.
; APPLICANT: DUGGAN, Brendan M.; LU, Yan
; TITLE OF INVENTION: RECEPTORS
; FILE REFERENCE: PF-0793 USN
; CURRENT APPLICATION NUMBER: US/10/311,623
; CURRENT FILING DATE: 2002-12-17
; PRIOR APPLICATION NUMBER: US 01/19942
; PRIOR FILING DATE: 2001-06-21
; PRIOR APPLICATION NUMBER: US 60/214,027
; PRIOR FILING DATE: 2000-06-21
; PRIOR APPLICATION NUMBER: US 60/228,045
; PRIOR FILING DATE: 2000-08-25
; PRIOR APPLICATION NUMBER: US 60/255,104
; PRIOR FILING DATE: 2000-12-12
; NUMBER OF SEQ ID NOS: 24
; SOFTWARE: PERL Program
; SEQ ID NO 13
;   LENGTH: 3580
;   TYPE: DNA
;   ORGANISM: Homo sapiens
; ..FEATURE:
;   NAME/KEY: misc_feature
;   OTHER INFORMATION: Incyte ID No. US20040023244A1 6052371CB1
US-10-311-623-13

  Query Match   86.9%; Score 2343; DB 17; Length 3580;
  Best Local Similarity 93.6%; Pred. No. 0;
  Matches 2524; Conservative   0; Mismatches   5; Indels 168; Gaps   1;

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Qy      1 ATGGCCGTCCGGCCCGGCCTGTGGCCAGCGCTCCTGGGCATAGTCCTCGCCGCTTGGCTC 60
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Db      4 ATGGCCGTCCGGCCCGGCCTGTGGCCAGCGCTCCTGGGCATAGTCCTCGCCGCTTGGCTC 63
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Qy    61 CGCGGCTCGGGTGCCCAGCAGAGTGCCACCGTGGCCAACCCAGTGCCTGGTGCCAACCCG 120
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Db    64 CGCGGCTCGGGTGCCCAGCAGAGTGCCACCGTGGCCAACCCAGTGCCTGGTGCCAACCCG 123

Qy   121 GACCTGCTTCCCCACTTCCTGGTGGAGCCCGAGGATGTGTACATCGTCAAGAACAAGCCA 180
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Qy   181 GTGCTGCTTGTGTGCAAGGCCGTGCCCGCCACGCAGATCTTCTTCAAGTGCAACGGGGAG 240
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Qy   241 TGGGTGCGCCAGGTGGACCACGTGATCGAGCGCAGCACAGACGGGAGCAGTGGGCTGCCC 300
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Qy   301 ACCATGGAGGTCCGCATTAATGTCTCAAGGCAGCAGGTCGAGAAGGTGTTCGGGCTGGAG 360
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Db   304 ACCATGGAGGTCCGCATTAATGTCTCAAGGCAGCAGGTCGAGAAGGTGTTCGGGCTGGAG 363

Qy   361 GAATACTGGTGCCAGTGCGTGGCATGGAGCTCCTCGGGCACCACCAAGAGTCAGAAGGCC 420
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Db   364 GAATACTGGTGCCAGTGCGTGGCATGGAGCTCCTCGGGCACCACCAAGAGTCAGAAGGCC 423

Qy   421 TACATCCGCATAGCCAGATTGCGCAAGAACTTCGAGCAGGAGCCGCTGGCCAAGGAGGTG 480
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Db   424 TACATCCGCATAGCCTATTTGCGCAAGAACTTCGAGCAGGAGCCGCTGGCCAAGGAGGTG 483

Qy   481 TCCCTGGAGCAGGGCATCGTGCTGCCCTGCCGTCCACCGGAGGGCATCCCTCCAGCCGAG 540
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Db   484 TCCCTGGAGCAGGGCATCGTGCTGCCCTGCCGTCCACCGGAGGGCATCCCTCCAGCCGAG 543

Qy   541 GTGGAGTGGCTCCGGAACGAGGACCTGGTGGACCCGTCCCTGGACCCCAATGTATACATC 600
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Qy   601 ACGCGGGAGCACAGCCTGGTGGTGCGACAGGCCCGCCTTGCTGACACGGCCAACTACACC 660
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Db   604 ACGCGGGAGCACAGCCTGGTGGTGCGACAGGCCCGCCTTGCTGACACGGCCAACTACACC 663

Qy   661 TGCGTGGCCAAGAACATCGTGGCACGTCGCCGCAGCGCCTCCGCTGCTGTCATCGTCTAC 720
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Db   664 TGCGTGGCCAAGAACATCGTGGCACGTCGCCGCAGCGCCTCCGCTGCTGTCATCGTCTAC 723

Qy   721 GTGAACGGTGGGTGGTCGACGTGGACCGAGTGGTCCGTCTGCAGCGCCAGCTGTGGGCGC 780
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Db   724 GTG--------------------------------------------------------- 726

Qy   781 GGCTGGCAGAAACGGAGCCGGAGCTGCACCAACCCGGCGCCTCTCAACGGGGGCGCTTTC 840

Db   727 --------------------------------------------------------- 726

Qy   841 TGTGAGGGGCAGAATGTCCAGAAAACAGCCTGCGCCACCCTGTGCCCAGTAGACGGCAGC 900
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Qy    901 TGGAGCCCGTGGAGCAAGTGGTCGGCCTGTGGGCTGGACTGCACCCACTGGCGGAGCCGT 960
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Db    736 TGGAGCCCGTGGAGCAAGTGGTCGGCCTGTGGGCTGGACTGCACCCACTGGCGGAGCCGT 795

Qy    961 GAGTGCTCTGACCCAGCACCCCGCAACGGAGGGGAGGAGTGCCAGGGCACTGACCTGGAC 1020
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    796 GAGTGCTCTGACCCAGCACCCCGCAACGGAGGGGAGGAGTGCCAGGGCACTGACCTGGAC 855

Qy   1021 ACCCGCAACTGTACCAGTGACCTCTGTGTACACAGTGCTTCTGGCCCTGAGGACGTGGCC 1080
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Db    856 ACCCGCAACTGTACCAGTGACCTCTGTGTACACACTGCTTCTGGCCCTGAGGACGTGGCC 915

Qy   1081 CTCTATGTGGGCCTCATCGCCGTGGCCGTCTGCCTGGTCCTGCTGCTGCTTGTCCTCATC 1140
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Db    916 CTCTATGTGGGCCTCATCGCCGTGGCCGTCTGCCTGGTCCTGCTGCTGCTTGTCCTCATC 975

Qy   1141 CTCGTTTATTGCCGGAAGAAGGAGGGGCTGGACTCAGATGTGGCTGACTCGTCCATTCTC 1200
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Db    976 CTCGTTTATTGCCGGAAGAAGGAGGGGCTGGACTCAGATGTGGCTGACTCGTCCATTCTC 1035

Qy   1201 ACCTCAGGCTTCCAGCCCGTCAGCATCAAGCCCAGCAAAGCAGACAACCCCCATCTGCTC 1260
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Db   1036 ACCTCAGGCTTCCAGCCCGTCAGCATCAAGCCCAGCAAAGCAGACAACCCCCATCTGCTC 1095

Qy   1261 ACCATCCAGCCGGACCTCAGCACCACCACCACCACCTACCAGGGCAGTCTCTGTCCCCGG 1320
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Db   1096 ACCATCCAGCCGGACCTCAGCACCACCACCACCACCTACCAGGGCAGTCTCTGTCCCCGG 1155

Qy   1321 CAGGATGGGCCCAGCCCCAAGTTCCAGCTCACCAATGGGCACCTGCTCAGCCCCCTGGGT 1380
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Db   1156 CAGGATGGGCCCAGCCCCAAGTTCCAGCTCACCAATGGGCACCTGCTCAGCCCCCTGGGT 1215

Qy   1381 GGCGGCCGCCACACACTGCACCACAGCTCTCCCACCTCTGAGGCCGAGGAGTTCGTCTCC 1440
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Db   1216 GGCGGCCGCCACACACTGCACCACAGCTCTCCCACCTCTGAGGCCGAGGAGTTCGTCTCC 1275

Qy   1441 CGCCTCTCCACCCAGAACTACTTCCGCTCCCTGCCCCGAGGCACCAGCAACATGACCTAT 1500
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Db   1276 CGCCTCTCCACCCAGAACTACTTCCGCTCCCTGCCCCGAGGCACCAGCAACATGACCTAT 1335

Qy   1501 GGGACCTTCAACTTCCTCGGGGGCCGGCTGATGATCCCTAATACAGGTATCAGCCTCCTC 1560
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Db   1396 ATCCCCCCAGATGCCATACCCCGAGGGAAGATCTATGAGATCTACCTCACGCTGCACAAG 1455

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Qy   1741 GAGCCCAGCCCTGACAGCTGGAGCCTGCGCCTCAAAAAGCAGTCGTGCGAGGGCAGCTGG 1800
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Db   1576 GAGCCCAGCCCTGACAGCTGGAGCCTGCGCCTCAAAAAGCAGTCGTGCGAGGGCAGCTGG 1635

Qy   1801 GAGGATGTGCTGCACCTGGGCGAGGAGGCGCCCTCCCACCTCTACTACTGCCAGCTGGAG 1860
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Db   1636 GAGGATGTGCTGCACCTGGGCGAGGAGGCGCCCTCCCACCTCTACTACTGCCAGCTGGAG 1695

Qy   1861 GCCAGTGCCTGCTACGTCTTCACCGAGCAGCTGGGCCGCTTTGCCCTGGTGGGAGAGGCC 1920
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Db   1816 TCCCTCGAGTACAACATCCGGGTCTACTGCCTGCATGACACCCACGATGCACTCAAGGAG 1875

Qy   2041 GTGGTGCAGCTGGAGAAGCAGCTGGGGGGGACAGCTGATCCAGGAGCCACGGGTCCTGCAC 2100
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Qy   2101 TTCAAGGACAGTTACCACAACCTGCGCCTATCCATCCACGATGTGCCCAGCTCCCTGTGG 2160
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Db   1936 TTCAAGGACAGTTACCACAACCTGCGCCTATCCATCCACGATGTGCCCAGCTCCCTGTGG 1995

Qy   2161 AAGAGTAAGCTCCTTGTCAGCTACCAGGAGATCCCCTTTTATCACATCTGGAATGGCACG 2220
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Db   1996 AAGAGTAAGCTCCTTGTCAGCTACCAGGAGATCCCCTTTTATCACATCTGGAATGGCACG 2055

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Db   2116 GCCTGCAAGCTGTGGGTGTGGCAGGTGGAGGGCGACGGGCAGAGCTTCAGCATCAACTTC 2175

Qy   2341 AACATCACCAAGGACACAAGGTTTGCTGAGCTGCTGGCTCTGGAGAGTGAAGCGGGGGTC 2400
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Db   2176 AACATCACCAAGGACACAAGGTTTGCTGAGCTGCTGGCTCTGGAGAGTGAAGCGGGGGTC 2235

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Db      2416 AACCTGTGGGAGGCGCGGCACTTCCCCAACGGCAACCTCAGCCAGCTGGCTGCAGCAGTG 2475

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Db      2476 GCTGGACTGGGCCAGCCAGACGCTGGCCTCTTCACAGTGTCGGAGGCTGAGTGCTGA 2532
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RESULT 6
US-10-240-154-15
; Sequence 15, Application US/10240154
; Publication No. US20030175741A1
; GENERAL INFORMATION:
; APPLICANT: Cochran et al.
; TITLE OF INVENTION: SCHIZOPHRENIA RELATED GENES
; FILE REFERENCE: CKFW-P01-006
; CURRENT APPLICATION NUMBER: US/10/240,154
; CURRENT FILING DATE: 2001-04-02
; PRIOR APPLICATION NUMBER: PCT/GB01/01486
; PRIOR FILING DATE: 2001-04-02
; NUMBER OF SEQ ID NOS: 34
; SOFTWARE: PatentIn version 3.2
; SEQ ID NO 15
;   LENGTH: 2697
;   TYPE: DNA
;   ORGANISM: Rattus sp.
;   FEATURE:
;   NAME/KEY: CDS
;   LOCATION: (1)..(2697)
US-10-240-154-15

  Query Match   83.5%; Score 2252.2; DB 16; Length 2697;
  Best Local Similarity   89.7%; Pred. No. 0;
  Matches 2419; Conservative   0; Mismatches 278; Indels   0; Gaps   0;

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Qy   61 CGCGGCTCGGGTGCCCAGCAGAGTGCCACCGTGGCCAACCCAGTGCCTGGTGCCAACCCG 120
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Db   61 CGTGGTTCGGGTGCCCAGCAGAGTGCCACGGTGGCCAATCCAGTGCCCGGTGCCAACCCC 120

Qy   121 GACCTGCTTCCCCACTTCCTGGTGGAGCCCGAGGATGTGTACATCGTCAAGAACAAGCCA 180
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Qy   181 GTGCTGCTTGTGTGCAAGGCCGTGCCCGCCACGCAGATCTTCTTCAAGTGCAACGGGGAG 240
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Db   181 GTGTTGTTGGTGTGCAAGGCTGTGCCTGCCACCCAGATCTTCTTCAAGTGCAATGGGGAA 240

Qy   241 TGGGTGCGCCAGGTGGACCACGTGATCGAGCGCAGCACAGACGGGAGCAGTGGGCTGCCC 300
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Db   241 TGGGTCCGCCAGGTCGATCACGTAATTGAACGCAGCACCGACAGCAGCAGCGGATTGCCA 300

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Db   301 ACCATGGAGGTCCGTATCAACGTATCGAGGCAGCAGGTAGAGAAAGTGTTTGGGCTGGAG 360
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Qy     361 GAATACTGGTGCCAGTGCGTGGCATGGAGCTCCTCGGGCACCACCAAGAGTCAGAAGGCC 420
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Db     361 GAATACTGGTGCCAGTGTGTGGCATGGAGCTCCTCGGGTACCACCAAAAGTCAGAAGGCC 420

Qy     421 TACATCCGCATAGCCAGATTGCGCAAGAACTTCGAGCAGGAGCCGCTGGCCAAGGAGGTG 480
           |||||||| || |||    ||||||||||||||| |||||||||||| |||||||||| |||
Db     421 TACATCCGGATTGCCTATTTGCGCAAGAACTTTGAGCAGGAGCCACTGGCCAAGGAAGTG 480

Qy     481 TCCCTGGAGCAGGGCATCGTGCTGCCCTGCCGTCCACCGGAGGGCATCCCTCCAGCCGAG 540
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Db     481 TCACTGGAGCAAGGCATTGTACTACCTTGTCGCCCCCCAGAAGGAATCCCCCCAGCTGAG 540

Qy     541 GTGGAGTGGCTCCGGAACGAGGACCTGGTGGACCCGTCCCTGGACCCCAATGTATACATC 600
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Db     541 GTGGAGTGGCTTCGAAATGAGGACCTCGTGGACCCCTCCCTCGATCCCAATGTGTACATC 600

Qy     601 ACGCGGGAGCACAGCCTGGTGGTGCGACAGGCCCGCCTTGCTGACACGGCCAACTACACC 660
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Db     601 ACGCGGGAGCACAGCCTAGTCGTGCGTCAGGCCCGCCTGGCCGACACGGCCAACTACACC 660

Qy     661 TGCGTGGCCAAGAACATCGTGGCACGTCGCCGCAGCGCCTCCGCTGCTGTCATCGTCTAC 720
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Db     661 TGTGTGGCCAAGAACATCGTAGCCCGTCGCCGAAGCACCTCTGCAGCGGTCATTGTTTAT 720

Qy     721 GTGAACGGTGGGTGGTCGACGTGGACCGAGTGGTCCGTCTGCAGCGCCAGCTGTGGGCGC 780
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Qy     781 GGCTGGCAGAAACGGAGCCGGAGCTGCACCAACCCGGCGCCTCTCAACGGGGGCGCTTTC 840
           |||||||||||||||||||||||||||||||||||||||| |||||||||||||||| |||
Db     781 GGCTGGCAGAAACGGAGCCGGAGCTGCACCAACCCGGCACCTCTCAACGGGGGCGCCTTC 840

Qy     841 TGTGAGGGGCAGAATGTCCAGAAAACAGCCTGCGCCACCCTGTGCCCAGTAGACGGCAGC 900
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Db     841 TGTGAGGGGCAGAATGTCCAGAAAACAGCCTGCGCCACTCTGTGCCCAGTGGATGGGAGC 900

Qy     901 TGGAGCCCGTGGAGCAAGTGGTCGGCCTGTGGGCTGGACTGCACCCACTGGCGGAGCCGT 960
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Db     901 TGGAGTTCGTGGAGTAAGTGGTCAGCCTGTGGGCTTGACTGCACCCACTGGCGGAGCCGC 960

Qy     961 GAGTGCTCTGACCCAGCACCCCGCAACGGAGGGGAGGAGTGCCAGGGCACTGACCTGGAC 1020
           |||||||||||||||||||||||||||| ||||| ||||||| | ||| ||||||||||||
Db     961 GAGTGCTCTGACCCAGCACCCCGCAATGGAGGTGAGGAGTGTCGGGGTGCTGACCTGGAC 1020

Qy    1021 ACCCGCAACTGTACCAGTGACCTCTGTGTACACAGTGCTTCTGGCCCTGAGGACGTGGCC 1080
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Db    1021 ACCCGCAACTGTACCAGTGACCTCTGCCTGCACACCGCTTCTTGCCCCGAGGACGTGGCT 1080

Qy    1081 CTCTATGTGGGCCTCATCGCCGTGGCCGTCTGCCTGGTCCTGCTGCTGCTTGTCCTCATC 1140
           |||||   | ||||||    |||| ||||| || |||||   || |||||| |||| | |||
Db    1081 CTCTACATCGGCCTTGTCGCTGTGGCTGTGTGCCTCTTCTTGCTGTTGCTGGCCCTTGGA 1140

Qy    1141 CTCGTTTATTGCCGGAAGAAGGAGGGGCTGGACTCAGATGTGGCTGACTCGTCCATTCTC 1200
           ||| |||| || || ||||||||| ||||||||||||| |||||||||||||||||| |||
Db    1141 CTCATTTACTGTCGCAAGAAGGAAGGGCTGGACTCCGATGTGGCCGACTCGTCCATCCTC 1200
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Qy    1201 ACCTCAGGCTTCCAGCCCGTCAGCATCAAGCCCAGCAAAGCAGACAACCCCCATCTGCTC 1260
           ||||| ||||||||||| |||||||||||||||||||||||||||||||||||| ||||||
Db    1201 ACCTCGGGCTTCCAGCCTGTCAGCATCAAGCCCAGCAAAGCAGACAACCCCCACCTGCTC 1260

Qy    1261 ACCATCCAGCCGGACCTCAGCACCACCACCACCACCTACCAGGGCAGTCTCTGTCCCCGG 1320
           |||||||||| |||||||||||||||||||||| |||||||||||||||| ||| | | ||
Db    1261 ACCATCCAGCCAGACCTCAGCACCACCACTACCACCTACCAGGGCAGTCTATGTTCGAGG 1320

Qy    1321 CAGGATGGGCCCAGCCCCAAGTTCCAGCTCACCAATGGGCACCTGCTCAGCCCCCTGGGT 1380
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Db    1321 CAGGATGGACCCAGCCCCAAGTTCCAGCTCTCTAATGGTCACCTGCTCAGCCCACTGGGG 1380

Qy    1381 GGCGGCCGCCACACACTGCACCACAGCTCTCCCACCTCTGAGGCCGAGGAGTTCGTCTCC 1440
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Db    1381 AGTGGCCGCCATACGTTGCACCACAGCTCACCCACCTCTGAGGCTGAGGACTTCGTCTCC 1440

Qy    1441 CGCCTCTCCACCCAGAACTACTTCCGCTCCCTGCCCCGAGGCACCAGCAACATGACCTAT 1500
           |||||||||||||| ||||||| || ||||||||||| ||||||||||||||| ||||
Db    1441 CGCCTCTCCACCCAAAACTACTTTCGTTCCCTGCCCCGCGGCACCAGCAACATGGCCTAC 1500

Qy    1501 GGGACCTTCAACTTCCTCGGGGGCCGGCTGATGATCCCTAATACAGGTATCAGCCTCCTC 1560
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Db    1501 GGGACCTTCAACTTCCTCGGGGGCCGGCTGATGATCCCTAATACGGGGATCAGCCTCCTC 1560

Qy    1561 ATCCCCCCAGATGCCATACCCCGAGGGAAGATCTATGAGATCTACCTCACGCTGCACAAG 1620
           || ||||| ||||||||| ||||||| ||||||||| |||||||||||||| ||||||||
Db    1561 ATACCCCCGGATGCCATCCCCCGAGGAAAGATCTACGAGATCTACCTCACACTGCACAAG 1620

Qy    1621 CCGGAAGACGTGAGGTTGCCCCTAGCTGGCTGTCAGACCCTGCTGAGTCCCATCGTTAGC 1680
           || |||||||||||||||||||||||||||||||||||||||||||||||| ||||||||
Db    1621 CCAGAAGACGTGAGGTTGCCCCTAGCTGGCTGTCAGACCCTGCTGAGTCCAGTCGTTAGC 1680

Qy    1681 TGTGGACCCCCTGGCGTCCTGCTCACCCGGCCAGTCATCCTGGCTATGGACCACTGTGGG 1740
           ||||| |||||| || ||||||||||||||||||||||||||| || |||||||||||||
Db    1681 TGTGGGCCCCCAGGAGTCCTGCTCACCCGGCCAGTCATCCTTGCAATGGACCACTGTGGA 1740

Qy    1741 GAGCCCAGCCCTGACAGCTGGAGCCTGCGCCTCAAAAAGCAGTCGTGCGAGGGCAGCTGG 1800
           ||||||||||||||||||||||| | ||||||||||||||||||| ||||||||||| |||
Db    1741 GAGCCCAGCCCTGACAGCTGGAGTCTGCGCCTCAAAAAGCAGTCCTGCGAGGGCAGTTGG 1800

Qy    1801 GAGGATGTGCTGCACCTGGGCGAGGAGGCGCCCTCCCACCTCTACTACTGCCAGCTGGAG 1860
           |||||||||||||||| || || |||||| | || ||||||||||||||||||||||||||
Db    1801 GAGGATGTGCTGCACCTTGGTGAGGAGTCACCTTCCCACCTCTACTACTGCCAGCTGGAG 1860

Qy    1861 GCCAGTGCCTGCTACGTCTTCACCGAGCAGCTGGGCCGCTTTGCCCTGGTGGGAGAGGCC 1920
           ||| | |||.|.||| ||||||||| |||||||||||||||||||||||||| ||||||||
Db    1861 GCCGGGGCCTGCTATGTCTTCACGGAGCAGCTGGGCCGCTTTGCCCTGGTAGGAGAGGCC 1920

Qy    1921 CTCAGCGTGGCTGCCGCCAAGCGCCTCAAGCTGCTTCTGTTTGCGCCGGTGGCCTGCACC 1980
           |||||||||||||| ||||||||||||| ||| ||||||||||| || |||||||| ||
Db    1921 CTCAGCGTGGCTGCCACCAAGCGCCTCAGGCTCCTTCTGTTTGCTCCCGTGGCCTGTACG 1980

Qy    1981 TCCCTCGAGTACAACATCCGGGTCTACTGCCTGCATGACACCCACGATGCACTCAAGGAG 2040
           ||||| |||||||||||||| || |||||||| || |||||||||||| || ||||||||
Db    1981 TCCCTTGAGTACAACATCCGAGTGTACTGCCTACACGACACCCACGACGCTCTCAAGGAG 2040

Qy    2041 GTGGTGCAGCTGGAGAAGCAGCTGGGGGGACAGCTGATCCAGGAGCCACGGGTCCTGCAC 2100
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                   ||||||||||||||||||||||| || |||||||||||||||||||||| || ||||||||||
Db    2041 GTGGTGCAGCTGGAGAAGCAGCTAGGTGGACAGCTGATCCAGGAGCCTCGCGTCCTGCAC 2100

Qy    2101 TTCAAGGACAGTTACCACAACCTGCGCCTATCCATCCACGATGTGCCCAGCTCCCTGTGG 2160
                   ||||| ||||||||||||||||| || || |||||||||||| ||||||||||||||||
Db    2101 TTCAAAGACAGTTACCACAACCTACGTCTCTCCATCCACGACGTGCCCAGCTCCCTGTGG 2160

Qy    2161 AAGAGTAAGCTCCTTGTCAGCTACCAGGAGATCCCCTTTTATCACATCTGGAATGGCACG 2220
                   ||||| |||||| |||||||||||||||||||||||| ||||| |||||||||| |||||
Db    2161 AAGAGCAAGCTACTTGTCAGCTACCAGGAGATCCCTTTTTACCACATCTGGAACGGCACC 2220

Qy    2221 CAGCGGTACTTGCACTGCACCTTCACCCTGGAGCGTGTCAGCCCCAGCACTAGTGACCTG 2280
                   |||| ||| |||||||||||||||||||||||||||| ||| | ||||||| || ||||||
Db    2221 CAGCAGTATCTGCACTGCACCTTCACCCTGGAGCGCATCAACGCCAGCACCAGCGACCTG 2280

Qy    2281 GCCTGCAAGCTGTGGGTGTGGCAGGTGGAGGGCGACGGGCAGAGCTTCAGCATCAACTTC 2340
                   ||||||||| |||||||||||||||||||||| || |||||||||||||| ||||||||||
Db    2281 GCCTGCAAGGTGTGGGTGTGGCAGGTGGAGGGAGATGGGCAGAGCTTCAACATCAACTTC 2340

Qy    2341 AACATCACCAAGGACACAAGGTTTGCTGAGCTGCTGGCTCTGGAGAGTGAAGCGGGGGTC 2400
                   ||||||| ||||||||||||||||||||||  || |||||||||||||||||| |||||||
Db    2341 AACATCACTAAGGACACAAGGTTTGCTGAATTGTTGGCTCTGGAGAGTGAAGGGGGGGTC 2400

Qy    2401 CCAGCCCTGGTGGGCCCCAGTGCCTTCAAGATCCCCTTCCTCATTCGGCAGAAGATAATT 2460
                   ||||||||||||||||||||||||||||||||||||||||||||||||||||| ||||| ||
Db    2401 CCAGCCCTGGTGGGCCCCAGTGCCTTCAAGATCCCCTTCCTCATTCGGCAAAAGATCATC 2460

Qy    2461 TCCAGCCTGGACCCACCCTGTAGGCGGGGTGCCGACTGGCGGACTCTGGCCCAGAAACTC 2520
                    |||| |||||||||||||| || ||||| |||||||||| | ||||| ||||||||||
Db    2461 GCCAGTCTGGACCCACCCTGCAGCCGGGGCGCCGACTGGAGAACTCTAGCCCAGAAACTT 2520

Qy    2521 CACCTGGACAGCCATCTCAGCTTCTTTGCCTCCAAGCCCAGCCCCACAGCCATGATCCTC 2580
                   ||||||||||||||||||| |||||||||||||||||||||||||| |||||||||||||||
Db    2521 CACCTGGACAGCCATCTTAGCTTCTTTGCCTCCAAGCCCAGCCCTACAGCCATGATCCTC 2580

Qy    2581 AACCTGTGGGAGGCGCGGCACTTCCCCAACGGCAACCTCAGCCAGCTGGCTGCAGCAGTG 2640
                   ||||| ||||||||| ||||||||||||||||||||||| ||||||||||| ||||| |||
Db    2581 AACCTATGGGAGGCACGGCACTTCCCCAACGGCAACCTCGGCCAGCTGGCAGCAGCTGTG 2640

Qy    2641 GCTGGACTGGGCCAGCCAGACGCTGGCCTCTTCACAGTGTCGGAGGCTGAGTGCTGA 2697
                   || ||||||||||| ||||| |||||||||||||||| |||||||||| ||||| |||
Db    2641 GCCGGACTGGGCCAACCAGATGCTGGCCTCTTCACGGTGTCGGAGGCCGAGTGTTGA 2697
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RESULT 7
US-09-933-261-1
; Sequence 1, Application US/09933261
; Publication No. US20030040046A1
;   GENERAL INFORMATION:
;   APPLICANT: Tessier-Lavigne, Marc
;   Leonardo, E. David
;   Hink, Lindsay
;   Masu, Masayuki
;   Kazuko, Keino-Masu
;   TITLE OF INVENTION: Netrin Receptors
;   NUMBER OF SEQUENCES: 8
;   CORRESPONDENCE ADDRESS:

;   INFORMATION FOR SEQ ID NO: 1:
;   SEQUENCE CHARACTERISTICS:
;   LENGTH: 3014 base pairs
;   TYPE: nucleic acid
;   STRANDEDNESS: double
;   TOPOLOGY: linear
;   MOLECULE TYPE: cDNA
;   SEQUENCE DESCRIPTION: SEQ ID NO: 1:
US-09-933-261-1

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  Query Match              83.5%;  Score 2252.2;  DB 10;  Length 3014;
  Best Local Similarity    89.7%;  Pred. No. 0;
  Matches 2419;  Conservative    0;  Mismatches  278;  Indels    0;  Gaps    0;

Qy        1 ATGGCCGTCCGGCCCGGCCTGTGGCCAGCGCTCCTGGGCATAGTCCTCGCCGCTTGGCTC 60
            |||||||||||||||||||||||||||||| |||||||||||||||||||||||| |||||
Db        1 ATGGCCGTCCGGCCCGGCCTGTGGCCAGTGCTCCTGGGCATAGTCCTCGCCGCCTGGCTT 60

Qy       61 CGCGGCTCGGGTGCCCAGCAGAGTGCCACCGTGGCCAACCCAGTGCCTGGTGCCAACCCG 120
            || || |||||||||||||||||||||||| ||||||||| ||||||||| |||||||||||
Db       61 CGTGGTTCGGGTGCCCAGCAGAGTGCCACGGTGGCCAATCCAGTGCCCGGTGCCAACCCC 120

Qy      121 GACCTGCTTCCCCACTTCCTGGTGGAGCCCGAGGATGTGTACATCGTCAAGAACAAGCCA 180
            |||||||| ||||||||||||||| |||||| ||||| ||||| ||||||||| |||||||||||
Db      121 GACCTGCTGCCCCACTTCCTGGTAGAGCCTGAGGACGTGTACATTGTCAAGAACAAGCCG 180

Qy      181 GTGCTGCTTGTGTGCAAGGCCGTGCCCGCCACGCAGATCTTCTTCAAGTGCAACGGGGAG 240
            ||| || | |||||||||||| ||||| ||||| |||||||||||||||||||||| |||||
Db      181 GTGTTGTTGGTGTGCAAGGCTGTGCCTGCCACCCAGATCTTCTTCAAGTGCAATGGGGAA 240

Qy      241 TGGGTGCGCCAGGTGGACCACGTGATCGAGCGCAGCACAGACGGGAGCAGTGGGCTGCCC 300
            ||||| ||||||||| || |||||| || || ||||||||| ||| | ||||| || ||||
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Db   241 TGGGTCCGCCAGGTCGATCACGTAATTGAACGCAGCACCGACAGCAGCAGCGGATTGCCA 300

Qy   301 ACCATGGAGGTCCGCATTAATGTCTCAAGGCAGCAGGTCGAGAAGGTGTTCGGGCTGGAG 360
         ||||||||||||||| || || || || |||||||||||| ||||| ||||| ||||||||
Db   301 ACCATGGAGGTCCGTATCAACGTATCGAGGCAGCAGGTAGAGAAAGTGTTTGGGCTGGAG 360

Qy   361 GAATACTGGTGCCAGTGCGTGGCATGGAGCTCCTCGGGCACCACCAAGAGTCAGAAGGCC 420
         |||||||||||||||| ||||||||||||||||||||| |||||||| ||||||||||||
Db   361 GAATACTGGTGCCAGTGTGTGGCATGGAGCTCCTCGGGTACCACCAAAAGTCAGAAGGCC 420

Qy   421 TACATCCGCATAGCCAGATTGCGCAAGAACTTCGAGCAGGAGCCGCTGGCCAAGGAGGTG 480
         |||||||| || |||       |||||||||||| |||||||||| |||||||||| |||
Db   421 TACATCCGGATTGCCTATTTGCGCAAGAACTTTGAGCAGGAGCCACTGGCCAAGGAAGTG 480

Qy   481 TCCCTGGAGCAGGGCATCGTGCTGCCCTGCCGTCCACCGGAGGGCATCCCTCCAGCCGAG 540
         || |||||||| |||||| || || || || || || || || || || ||||| ||||| |||
Db   481 TCACTGGAGCAAGGCATTGTACTACCTTGTCGCCCCCCAGAAGGAATCCCCCCAGCTGAG 540

Qy   541 GTGGAGTGGCTCCGGAACGAGGACCTGGTGGACCCGTCCCTGGACCCCAATGTATACATC 600
         |||||||||| || || ||||||||| |||||||| ||||| || ||||||||| ||||||
Db   541 GTGGAGTGGCTTCGAAATGAGGACCTCGTGGACCCCTCCCTCGATCCCAATGTGTACATC 600

Qy   601 ACGCGGGAGCACAGCCTGGTGGTGCGACAGGCCCGCCTTGCTGACACGGCCAACTACACC 660
         ||||||||||||||||| || ||||| |||||||||||| || ||||||||||||||||||
Db   601 ACGCGGGAGCACAGCCTAGTCGTGCGTCAGGCCCGCCTGGCCGACACGGCCAACTACACC 660

Qy   661 TGCGTGGCCAAGAACATCGTGGCACGTCGCCGCAGCGCCTCCGCTGCTGTCATCGTCTAC 720
         || ||||||||||||||||||| || |||||| ||| |||| || || ||||| || ||
Db   661 TGTGTGGCCAAGAACATCGTAGCCCGTCGCCGAAGCACCTCTGCAGCGGTCATTGTTTAT 720

Qy   721 GTGAACGGTGGGTGGTCGACGTGGACCGAGTGGTCCGTCTGCAGCGCCAGCTGTGGGCGC 780
         |||||||||||||||||||||||||||| |||||||||||||||||||||||||||||||
Db   721 GTGAACGGTGGGTGGTCGACGTGGACTGAGTGGTCCGTCTGCAGCGCCAGCTGTGGGCGT 780

Qy   781 GGCTGGCAGAAACGGAGCCGGAGCTGCACCAACCCGGCGCCTCTCAACGGGGGCGCTTTC 840
         |||||||||||||||||||||||||||||||||||||||| ||||||||||||||| |||
Db   781 GGCTGGCAGAAACGGAGCCGGAGCTGCACCAACCCGGCACCTCTCAACGGGGGCGCCTTC 840

Qy   841 TGTGAGGGGCAGAATGTCCAGAAAACAGCCTGCGCCACCCTGTGCCCAGTAGACGGCAGC 900
         ||||||||||||||||||||||||||||||||||||||| |||||||||| || || |||
Db   841 TGTGAGGGGCAGAATGTCCAGAAAACAGCCTGCGCCACTCTGTGCCCAGTGGATGGGAGC 900

Qy   901 TGGAGCCCGTGGAGCAAGTGGTCGGCCTGTGGGCTGGACTGCACCCACTGGCGGAGCCGT 960
         |||||   ||||||| ||||||| ||||||||| ||||||||||||||||||||||||||
Db   901 TGGAGTTCGTGGAGTAAGTGGTCAGCCTGTGGGCTTGACTGCACCCACTGGCGGAGCCGC 960

Qy   961 GAGTGCTCTGACCCAGCACCCCGCAACGGAGGGGAGGAGTGCCAGGGCACTGACCTGGAC 1020
         ||||||||||||||||||||||||||| |||||| ||||||||| | |||   ||||||||||
Db   961 GAGTGCTCTGACCCAGCACCCCGCAATGGAGGTGAGGAGTGTCGGGGTGCTGACCTGGAC 1020

Qy  1021 ACCCGCAACTGTACCAGTGACCTCTGTGTACACAGTGCTTCTGGCCCTGAGGACGTGGCC 1080
         ||||||||||||||||||||||||||||| | |||| |||||| |||| |||||||||||
Db  1021 ACCCGCAACTGTACCAGTGACCTCTGCCTGCACACCGCTTCTTGCCCCGAGGACGTGGCT 1080

Qy  1081 CTCTATGTGGGCCTCATCGCCGTGGCCGTCTGCCTGGTCCTGCTGCTGCTTGTCCTCATC 1140
         |||||   |  ||||||   |||| ||||| || ||||||   || |||||| |||| | |||
Db  1081 CTCTACATCGGCCTTGTCGCTGTGGCTGTGTGCCTCTTCTTGCTGTTGCTGGCCCTTGGA 1140
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Qy    1141 CTCGTTTATTGCCGGAAGAAGGAGGGGCTGGACTCAGATGTGGCTGACTCGTCCATTCTC 1200
           ||| |||| || || ||||||||| ||||||||||| ||||||| |||||||||| |||
Db    1141 CTCATTTACTGTCGCAAGAAGGAAGGGCTGGACTCCGATGTGGCCGACTCGTCCATCCTC 1200

Qy    1201 ACCTCAGGCTTCCAGCCCGTCAGCATCAAGCCCAGCAAAGCAGACAACCCCCATCTGCTC 1260
           ||||| |||||||||| |||||||||||||||||||||||||||||||||||| ||||||
Db    1201 ACCTCGGGCTTCCAGCCTGTCAGCATCAAGCCCAGCAAAGCAGACAACCCCCACCTGCTC 1260

Qy    1261 ACCATCCAGCCGGACCTCAGCACCACCACCACCACCTACCAGGGCAGTCTCTGTCCCCGG 1320
           |||||||||| ||||||||||||||||||| |||||||||||||||||| ||| | ||
Db    1261 ACCATCCAGCCAGACCTCAGCACCACCACTACCACCTACCAGGGCAGTCTATGTTCGAGG 1320

Qy    1321 CAGGATGGGCCCAGCCCCAAGTTCCAGCTCACCAATGGGCACCTGCTCAGCCCCCTGGGT 1380
           |||||||| |||||||||||||||||||| | |||||| |||||||||||||| |||||
Db    1321 CAGGATGGACCCAGCCCCAAGTTCCAGCTCTCTAATGGTCACCTGCTCAGCCCACTGGGG 1380

Qy    1381 GGCGGCCGCCACACACTGCACCACAGCTCTCCCACCTCTGAGGCCGAGGAGTTCGTCTCC 1440
           | ||||||||| || ||||||||||||||| ||||||||||||| ||||| |||||||||
Db    1381 AGTGGCCGCCATACGTTGCACCACAGCTCACCCACCTCTGAGGCTGAGGACTTCGTCTCC 1440

Qy    1441 CGCCTCTCCACCCAGAACTACTTCCGCTCCCTGCCCCGAGGCACCAGCAACATGACCTAT 1500
           ||||||||||||||| ||||||| || ||||||||||||| ||||||||||||| ||||
Db    1441 CGCCTCTCCACCCAAAACTACTTTCGTTCCCTGCCCCGCGGCACCAGCAACATGGCCTAC 1500

Qy    1501 GGGACCTTCAACTTCCTCGGGGGCCGGCTGATGATCCCTAATACAGGTATCAGCCTCCTC 1560
           |||||||||||||||||||||||||||||||||||||||||||||| || |||||||||
Db    1501 GGGACCTTCAACTTCCTCGGGGGCCGGCTGATGATCCCTAATACGGGGATCAGCCTCCTC 1560

Qy    1561 ATCCCCCCAGATGCCATACCCCGAGGGAAGATCTATGAGATCTACCTCACGCTGCACAAG 1620
           || ||||| ||||||||| |||||||| ||||||| |||||||||||||||| |||||||
Db    1561 ATACCCCCGGATGCCATCCCCCGAGGAAAGATCTACGAGATCTACCTCACACTGCACAAG 1620

Qy    1621 CCGGAAGACGTGAGGTTGCCCCTAGCTGGCTGTCAGACCCTGCTGAGTCCCATCGTTAGC 1680
           || ||||||||||||||||||||||||||||||||||||||||||||||||   ||||||||
Db    1621 CCAGAAGACGTGAGGTTGCCCCTAGCTGGCTGTCAGACCCTGCTGAGTCCAGTCGTTAGC 1680

Qy    1681 TGTGGACCCCCTGGCGTCCTGCTCACCCGGCCAGTCATCCTGGCTATGGACCACTGTGGG 1740
           ||||| |||||| || ||||||||||||||||||||||||||| || |||||||||||||
Db    1681 TGTGGGCCCCCAGGAGTCCTGCTCACCCGGCCAGTCATCCTTGCAATGGACCACTGTGGA 1740

Qy    1741 GAGCCCAGCCCTGACAGCTGGAGCCTGCGCCTCAAAAAGCAGTCGTGCGAGGGCAGCTGG 1800
           |||||||||||||||||||||||| ||||||||||||||||||||| ||||||||| |||
Db    1741 GAGCCCAGCCCTGACAGCTGGAGTCTGCGCCTCAAAAAGCAGTCCTGCGAGGGCAGTTGG 1800

Qy    1801 GAGGATGTGCTGCACCTGGGCGAGGAGGCGCCCTCCCACCTCTACTACTGCCAGCTGGAG 1860
           ||||||||||||||||| || |||||| | || |||||||||||||||||||||||||||
Db    1801 GAGGATGTGCTGCACCTTGGTGAGGAGTCACCTTCCCACCTCTACTACTGCCAGCTGGAG 1860

Qy    1861 GCCAGTGCCTGCTACGTCTTCACCGAGCAGCTGGGCCGCTTTGCCCTGGTGGGAGAGGCC 1920
           ||| | ||||||||| ||||||| ||||||||||||||||||||||||||| |||||||||
Db    1861 GCCGGGGCCTGCTATGTCTTCACGGAGCAGCTGGGCCGCTTTGCCCTGGTAGGAGAGGCC 1920

Qy    1921 CTCAGCGTGGCTGCCGCCAAGCGCCTCAAGCTGCTTCTGTTTGCGCCGGTGGCCTGCACC 1980
           |||||||||||||||||||||||||||||| ||| ||||||||||| || |||||||| ||
Db    1921 CTCAGCGTGGCTGCCACCAAGCGCCTCAGGCTCCTTCTGTTTGCTCCCGTGGCCTGTACG 1980
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Qy    1981  TCCCTCGAGTACAACATCCGGGTCTACTGCCTGCATGACACCCACGATGCACTCAAGGAG 2040
            |||||  |||||||||||||  ||  |||||||  ||  |||||.|||||  || ||.|||||||
Db    1981  TCCCTTGAGTACAACATCCGAGTGTACTGCCTACACGACACCCACGACGCTCTCAAGGAG 2040

Qy    2041  GTGGTGCAGCTGGAGAAGCAGCTGGGGGGACAGCTGATCCAGGAGCCACGGGTCCTGCAC 2100
            ||||||||||||||||||||||||||||  ||  |||||||||||||||||||  |||||||||
Db    2041  GTGGTGCAGCTGGAGAAGCAGCTAGGTGGACAGCTGATCCAGGAGCCTCGCGTCCTGCAC 2100

Qy    2101  TTCAAGGACAGTTACCACAACCTGCGCCTATCCATCCACGATGTGCCCAGCTCCCTGTGG 2160
            |||||  ||||||||||||||||||  ||  ||  ||||||||||  |||||||||||.||||||||
Db    2101  TTCAAAGACAGTTACCACAACCTACGTCTCTCCATCCACGACGTGCCCAGCTCCCTGTGG 2160

Qy    2161  AAGAGTAAGCTCCTTGTCAGCTACCAGGAGATCCCCTTTTATCACATCTGGAATGGCACG 2220
            |||||  |||||  ||||||||||||||||||||||||||  |||||  ||||||||||||  |||||
Db    2161  AAGAGCAAGCTACTTGTCAGCTACCAGGAGATCCCTTTTTACCACATCTGGAACGGCACC 2220

Qy    2221  CAGCGGTACTTGCACTGCACCTTCACCCTGGAGCGTGTCAGCCCCAGCACTAGTGACCTG 2280
            ||||  |||   '|||||||||||||||||||||||||||||||  |||  | |||||||  || ||||||
Db    2221  CAGCAGTATCTGCACTGCACCTTCACCCTGGAGCGCATCAACGCCAGCACCAGCGACCTG 2280

Qy    2281  GCCTGCAAGCTGTGGGTGTGGCAGGTGGAGGGCGACGGGCAGAGCTTCAGCATCAACTTC 2340
            ||||||||||  |||||||||||||||||||||||||  ||  |||||||||||||  ||||||||||
Db    2281  GCCTGCAAGGTGTGGGTGTGGCAGGTGGAGGGAGATGGGCAGAGCTTCAACATCAACTTC 2340

Qy    2341  AACATCACCAAGGACACAAGGTTTGCTGAGCTGCTGGCTCTGGAGAGTGAAGCGGGGGTC 2400
            ||||||||  |||||||||||||||||||||  ||  ||||||||||||||||||||  |||||||
Db    2341  AACATCACTAAGGACACAAGGTTTGCTGAATTGTTGGCTCTGGAGAGTGAAGGGGGGGTC 2400

Qy    2401  CCAGCCCTGGTGGGCCCCAGTGCCTTCAAGATCCCCTTCCTCATTCGGCAGAAGATAATT 2460
            |||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||  ||||| ||
Db    2401  CCAGCCCTGGTGGGCCCCAGTGCCTTCAAGATCCCCTTCCTCATTCGGCAAAAGATCATC 2460

Qy    2461  TCCAGCCTGGACCCACCCTGTAGGCGGGGTGCCGACTGGCGGACTCTGGCCCAGAAACTC 2520
            ||||  ||||||||||||||||  ||  |||||  |||||||||  | |||||  |||||||||||
Db    2461  GCCAGTCTGGACCCACCCTGCAGCCGGGGCGCCGACTGGAGAACTCTAGCCCAGAAACTT 2520

Qy    2521  CACCTGGACAGCCATCTCAGCTTCTTTGCCTCCAAGCCCAGCCCCACAGCCATGATCCTC 2580
            |||||||||||||||||||||  ||||||||||||||||||||||||||||  ||||||||||||||||
Db    2521  CACCTGGACAGCCATCTTAGCTTCTTTGCCTCCAAGCCCAGCCCTACAGCCATGATCCTC 2580

Qy    2581  AACCTGTGGGAGGCGCGGCACTTCCCCAACGGCAACCTCAGCCAGCTGGCTGCAGCAGTG 2640
            |||||  ||||||||  |||||||||||||||||||||||||||||||||||  |||||||  ||||| |||
Db    2581  AACCTATGGGAGGCACGGCACTTCCCCAACGGCAACCTCGGCCAGCTGGCAGCAGCTGTG 2640

Qy    2641  GCTGGACTGGGCCAGCCAGACGCTGGCCTCTTCACAGTGTCGGAGGCTGAGTGCTGA 2697
            ||  |||||||||||  ||.|||  |||||||||||||||||||  ||||||||||||  |||  |||
Db    2641  GCCGGACTGGGCCAACCAGATGCTGGCCTCTTCACGGTGTCGGAGGCCGAGTGTTGA 2697
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RESULT 8
US-10-256-702-1
; Sequence 1, Application US/10256702
; Publication No. US20030059859A1
;   GENERAL INFORMATION:
;   APPLICANT: Tessier-Lavigne, Marc
;   Leonardo, E. David
;   Hink, Lindsay

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  Query Match          83.5%;  Score 2252.2;  DB 14;  Length 3014;
  Best Local Similarity  89.7%;  Pred. No. 0;
  Matches 2419;  Conservative   0;  Mismatches  278;  Indels   0;  Gaps    0;

Qy        1 ATGGCCGTCCGGCCCGGCCTGTGGCCAGCGCTCCTGGGCATAGTCCTCGCCGCTTGGCTC 60
            ||||||||||||||||||||||||||||| |||||||||||||||||||||||||| |||||
Db        1 ATGGCCGTCCGGCCCGGCCTGTGGCCAGTGCTCCTGGGCATAGTCCTCGCCGCCTGGCTT 60

Qy       61 CGCGGCTCGGGTGCCCAGCAGAGTGCCACCGTGGCCAACCCAGTGCCTGGTGCCAACCCG 120
            || ||  |||||||||||||||||||||||| ||||||||| |||||||| ||||||||||
Db       61 CGTGGTTCGGGTGCCCAGCAGAGTGCCACGGTGGCCAATCCAGTGCCCGGTGCCAACCCC 120

Qy      121 GACCTGCTTCCCCACTTCCTGGTGGAGCCCGAGGATGTGTACATCGTCAAGAACAAGCCA 180
            |||||||| |||||||||||||| ||||| ||||| ||||||||| ||||||||||||||||
Db      121 GACCTGCTGCCCCACTTCCTGGTAGAGCCTGAGGACGTGTACATTGTCAAGAACAAGCCG 180
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Qy   181 GTGCTGCTTGTGTGCAAGGCCGTGCCCGCCACGCAGATCTTCTTCAAGTGCAACGGGGAG 240
         ||| || | ||||||||||| ||||| ||||| ||||||||||||||||||||| |||||
Db   181 GTGTTGTTGGTGTGCAAGGCTGTGCCTGCCACCCAGATCTTCTTCAAGTGCAATGGGGAA 240

Qy   241 TGGGTGCGCCAGGTGGACCACGTGATCGAGCGCAGCACAGACGGGAGCAGTGGGCTGCCC 300
         ||||| |||||||| || |||||| || || ||||||||| ||| | |||||| ||  ||||
Db   241 TGGGTCCGCCAGGTCGATCACGTAATTGAACGCAGCACCGACAGCAGCAGCGGATTGCCA 300

Qy   301 ACCATGGAGGTCCGCATTAATGTCTCAAGGCAGCAGGTCGAGAAGGTGTTCGGGCTGGAG 360
         ||||||||||||| || || || || |||||||||| ||||| ||||| |||||||||
Db   301 ACCATGGAGGTCCGTATCAACGTATCGAGGCAGCAGGTAGAGAAAGTGTTTGGGCTGGAG 360

Qy   361 GAATACTGGTGCCAGTGCGTGGCATGGAGCTCCTCGGGCACCACCAAGAGTCAGAAGGCC 420
         |||||||||||||||||| |||||||||||||||||||| |||||||| ||||||||||||
Db   361 GAATACTGGTGCCAGTGTGTGGCATGGAGCTCCTCGGGTACCACCAAAAGTCAGAAGGCC 420

Qy   421 TACATCCGCATAGCCAGATTGCGCAAGAACTTCGAGCAGGAGCCGCTGGCCAAGGAGGTG 480
         |||||||| || ||| |||||||||||||||| |||||||||||| |||||||||||| |||
Db   421 TACATCCGGATTGCCTATTTGCGCAAGAACTTTGAGCAGGAGCCACTGGCCAAGGAAGTG 480

Qy   481 TCCCTGGAGCAGGGCATCGTGCTGCCCTGCCGTCCACCGGAGGGCATCCCTCCAGCCGAG 540
         || ||||||||| ||||| || || || || || || || || || || ||||| ||||| |||
Db   481 TCACTGGAGCAAGGCATTGTACTACCTTGTCGCCCCCCAGAAGGAATCCCCCCAGCTGAG 540

Qy   541 GTGGAGTGGCTCCGGAACGAGGACCTGGTGGACCCGTCCCTGGACCCCAATGTATACATC 600
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Qy   601 ACGCGGGAGCACAGCCTGGTGGTGCGACAGGCCCGCCTTGCTGACACGGCCAACTACACC 660
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Qy   661 TGCGTGGCCAAGAACATCGTGGCACGTCGCCGCAGCGCCTCCGCTGCTGTCATCGTCTAC 720
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Qy   721 GTGAACGGTGGGTGGTCGACGTGGACCGAGTGGTCCGTCTGCAGCGCCAGCTGTGGGCGC 780
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Db   721 GTGAACGGTGGGTGGTCGACGTGGACTGAGTGGTCCGTCTGCAGCGCCAGCTGTGGGCGT 780

Qy   781 GGCTGGCAGAAACGGAGCCGGAGCTGCACCAACCCGGCGCCTCTCAACGGGGGCGCTTTC 840
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Db   781 GGCTGGCAGAAACGGAGCCGGAGCTGCACCAACCCGGCACCTCTCAACGGGGGCGCCTTC 840

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Qy   901 TGGAGCCCGTGGAGCAAGTGGTCGGCCTGTGGGCTGGACTGCACCCACTGGCGGAGCCGT 960
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Db   901 TGGAGTTCGTGGAGTAAGTGGTCAGCCTGTGGGCTTGACTGCACCCACTGGCGGAGCCGC 960

Qy   961 GAGTGCTCTGACCCAGCACCCCGCAACGGAGGGGAGGAGTGCCAGGGCACTGACCTGGAC 1020
         ||||||||||||||||||||||||||||| ||||| ||||||||| | ||| ||||||||||||
Db   961 GAGTGCTCTGACCCAGCACCCCGCAATGGAGGTGAGGAGTGTCGGGGTGCTGACCTGGAC 1020
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Qy   1021 ACCCGCAACTGTACCAGTGACCTCTGTGTACACAGTGCTTCTGGCCCTGAGGACGTGGCC 1080
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Db   1021 ACCCGCAACTGTACCAGTGACCTCTGCCTGCACACCGCTTCTTGCCCCGAGGACGTGGCT 1080

Qy   1081 CTCTATGTGGGCCTCATCGCCGTGGCCGTCTGCCTGGTCCTGCTGCTGCTTGTCCTCATC 1140
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Db   1081 CTCTACATCGGCCTTGTCGCTGTGGCTGTGTGCCTCTTCTTGCTGTTGCTGGCCCTTGGA 1140

Qy   1141 CTCGTTTATTGCCGGAAGAAGGAGGGGCTGGACTCAGATGTGGCTGACTCGTCCATTCTC 1200
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Db   1141 CTCATTTACTGTCGCAAGAAGGAAGGGCTGGACTCCGATGTGGCCGACTCGTCCATCCTC 1200

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Db   1201 ACCTCGGGCTTCCAGCCTGTCAGCATCAAGCCCAGCAAAGCAGACAACCCCCACCTGCTC 1260

Qy   1261 ACCATCCAGCCGGACCTCAGCACCACCACCACCACCTACCAGGGCAGTCTCTGTCCCCGG 1320
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Qy   1381 GGCGGCCGCCACACACTGCACCACAGCTCTCCCACCTCTGAGGCCGAGGAGTTCGTCTCC 1440
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Qy         1981 TCCCTCGAGTACAACATCCGGGTCTACTGCCTGCATGACACCCACGATGCACTCAAGGAG 2040
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Db         1981 TCCCTTGAGTACAACATCCGAGTGTACTGCCTACACGACACCCACGACGCTCTCAAGGAG 2040

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Qy         2101 TTCAAGGACAGTTACCACAACCTGCGCCTATCCATCCACGATGTGCCCAGCTCCCTGTGG 2160
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Db         2161 AAGAGCAAGCTACTTGTCAGCTACCAGGAGATCCCTTTTTACCACATCTGGAACGGCACC 2220

Qy         2221 CAGCGGTACTTGCACTGCACCTTCACCCTGGAGCGTGTCAGCCCCAGCACTAGTGACCTG 2280
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Qy         2281 GCCTGCAAGCTGTGGGTGTGGCAGGTGGAGGGCGACGGGCAGAGCTTCAGCATCAACTTC 2340
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Db         2281 GCCTGCAAGGTGTGGGTGTGGCAGGTGGAGGGAGATGGGCAGAGCTTCAACATCAACTTC 2340

Qy         2341 AACATCACCAAGGACACAAGGTTTGCTGAGCTGCTGGCTCTGGAGAGTGAAGCGGGGGTC 2400
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Db         2341 AACATCACTAAGGACACAAGGTTTGCTGAATTGTTGGCTCTGGAGAGTGAAGGGGGGGTC 2400

Qy         2401 CCAGCCCTGGTGGGCCCCAGTGCCTTCAAGATCCCCTTCCTCATTCGGCAGAAGATAATT 2460
                ||||||||||||||||||||||||||||||||||||||||||||||||||||||||  |||||  ||
Db         2401 CCAGCCCTGGTGGGCCCCAGTGCCTTCAAGATCCCCTTCCTCATTCGGCAAAAGATCATC 2460

Qy         2461 TCCAGCCTGGACCCACCCTGTAGGCGGGGTGCCGACTGGCGGACTCTGGCCCAGAAACTC 2520
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Db         2461 GCCAGTCTGGACCCACCCTGCAGCCGGGGCGCCGACTGGAGAACTCTAGCCCAGAAACTT 2520

Qy         2521 CACCTGGACAGCCATCTCAGCTTCTTTGCCTCCAAGCCCAGCCCCACAGCCATGATCCTC 2580
                |||||||||||||||||||  |||||||||||||||||||||||||||||  |||||||||||||||||
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RESULT 9

; Sequence 2, Application US/09933261 .
; Publication No. US20030040046A1
; GENERAL INFORMATION:
; APPLICANT: Tessier-Lavigne, Marc
; Leonardo, E. David
; Hink, Lindsay
; Masu, Masayuki
; Kazuko, Keino-Masu
; TITLE OF INVENTION: Netrin Receptors
; NUMBER OF SEQUENCES: 8
; CORRESPONDENCE ADDRESS:
; ADDRESSEE: SCIENCE & TECHNOLOGY LAW GROUP
; STREET: 268 BUSH STREET, SUITE 3200
; CITY: SAN FRANCISCO
; STATE: CALIFORNIA
; COUNTRY: USA
; ZIP: 94104
; COMPUTER READABLE FORM:
; MEDIUM TYPE: Floppy disk
; COMPUTER: IBM PC compatible
; OPERATING SYSTEM: PC-DOS/MS-DOS
; SOFTWARE: PatentIn Release #1.0, Version #1.30
; CURRENT APPLICATION DATA:
; APPLICATION NUMBER: US/09/933,261
; FILING DATE: 20-Aug-2001
; CLASSIFICATION: <Unknown>
; PRIOR APPLICATION DATA:
; APPLICATION NUMBER: 08/808,982
; FILING DATE: <Unknown>
; ATTORNEY/AGENT INFORMATION:
; NAME: OSMAN, RICHARD A
; REGISTRATION NUMBER: 36,627
; REFERENCE/DOCKET NUMBER: UC96-217
; TELECOMMUNICATION INFORMATION:
; TELEPHONE: (415) 343-4341
; TELEFAX: (415) 343-4342
; INFORMATION FOR SEQ ID NO: 2:
; SEQUENCE CHARACTERISTICS:
; LENGTH: 1787 base pairs
; TYPE: nucleic acid
; STRANDEDNESS: double
; TOPOLOGY: linear
; MOLECULE TYPE: cDNA
; SEQUENCE DESCRIPTION: SEQ ID NO: 2:
US-09-933-261-2

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   Query Match            57.6%;   Score 1552.4;  DB 10;   Length 1787;
   Best Local Similarity  98.5%;   Pred. No. 0;
   Matches 1651;  Conservative   0;  Mismatches   16;  Indels   9;  Gaps    8;

Qy      1025 GCAACTGTACCAGTGACCTCTGTGTACACAGTGCTTCTGGCCCTGAGGACGTGGCCCTCT 1084
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Db         1 GCAACTGTACCAGTGACCTCTG-GTACACACTGCTTCTGGCCCTGAGGACGTGGCCCTCT 59


Qy      1085 ATGTGGGCCTCATCGCCGTGGCCGTCTGCCTGGTCCTGCTGCTGCTTGTCCTCATCCTCG 1144
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Db     60 ATGTGGGCCTCATCGCCGTGGCCGTCTGCCTGGTCCTGCTGCTGCTTGTCCTCATCCTCG 119

Qy   1145 TTTATTGCCGGAAGAAGGAGGGGCTGGACTCAGATGTGGCTGACTCGTCCATTCTCACCT 1204
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Db    120 TTTATTGCCGGAAGAAGGAGGGGCTGGACTCAGATGTGGCTGACTCGTCCATTCTCACCT 179

Qy   1205 CAGGCTTCCAGCCCGTCAGCATC-AAGCCCAGCAAAGCAGACAACCCCCATCTGCTCACC 1263
          |||||||||||||||||||||||| |||||||||||||||||||||||||||||||||||
Db    180 CAGGCTTCCAGCCCGTCAGCATCTAAGCCCAGCAAAGCAGACAACCCCCATCTGCTCACC 239

Qy   1264 ATCCAGCCGGACCTCAGCACCACCACCACCACCTACCAGGGCAGTCTCTGTCCCCGGCAG 1323
          |||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    240 ATCCAGCCGGACCTCAGCACCACCACCACCACCTACCAGGGCAGTCTCTGTCCCCGGCAG 299

Qy   1324 GATGGGCCCAGCCCCAAGTTCCAGCTCACCAATGGGCACCTGCTCAGCCCCCTGGGTGGC 1383
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Db    300 GATGGGCCCAGCCCCAAGTTCCAGCTCACCAATGGGCACCTGCTCAGCCCCCTGGGTGGC 359

Qy   1384 GGCCGCCACACACTGCACCACAGCTCTCCCACCTCTGAGGCCGAGGAGTTCGTCTCCCGC 1443
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Db    360 GGCCGCCACACACTGCACCACAGCTCTCCCACCTCTGAGGCCGAGGAGTTCGTCTCCCGC 419

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Qy   1504 ACCTTCAACTTCCTCGGGGGCCGGCTGATGATCCCTAATACAGGTATCAGCCTCCTCATC 1563
          |||||||||||||||||||||||||||||||||||||||||||||||| |||||||||||
Db    480 ACCTTCAACTTCCTCGGGGGCCGGCTGATGATCCCTAATACAGGAATCAGCCTCCTCATC 539

Qy   1564 CCCCCAGATGCCATACCCCGAGGGAAGATCTATGAGATCTACCTCACGCTGCACAAGCCG 1623
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Db    540 CCCCCAGATGCCATACCCCGAGGGAAGATCTATGAGATCTACCTCACGCTGCACAAGCCG 599

Qy   1624 GAAGACGTGAGGTTGCCCCTAGCTGGCTGTCAGACCCTGCTGAGTCCCATCGTTAGCTGT 1683
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Db    600 GAAGACGTGAGGTTGCCCCTAGCTGGCTGTCAGACCCTGCTGAGTCCCATCGTTAGCTGT 659

Qy   1684 GGACCCCCTGGCGTCCTGCTCACCCGGCCAGTCATCCTGGCTATGGACCACTGTGGGGAG 1743
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Db    660 GGACCCCCTGGCGTCCTGCTCACCCGGCCAGTCATCCTGGCTATGGACCACTGTGGGGAG 719

Qy   1744 CCCAGCCCTGACAGCTGGAGCCTGCGCCTCAAAAAGCAGTCGTGCGAGGGCAGCTGGGAG 1803
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Db    720 CCCAGCCCTGACAGCTGGAGCCTGGCCCTCAAAAAGCAGTCGTGCGAGGG-AGCTGGGAG 778

Qy   1804 GATGTGCTGCACCTGGGCGAGGAGGCGCCCTCCCACCTCTACTACTGCCAGCTGGAGGCC 1863
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Db    779 GATGT-CTGCACCTGGGCGAGGAGGCGCCCTCCCACCTCTACTACTGCCAGCTGGAGGCC 837

Qy   1864 AGTGCCTGCTACGTCTTCACCGAGCAGCTGGGCCGCTTTGCCCTGGTGGGAGAGGCCCTC 1923
          |||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    838 AGTGCCTGCTACGTCTTCACCGAGCAGCTGGGCCGCTTTGCCCTGGTGGGAGAGGCCCTC 897

Qy   1924 AGCGTGGCTGCCGCCAAGCGCCTCAAGCTGCTTCTGTTTGCGCCGGTGGCCTGCACCTCC 1983
          |||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    898 AGCGTGGCTGCCGCCAAGCGCCTCAAGCTGCTTCTGTTTGCGCCGGTGGCCTGCACCTCC 957
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Qy    1984 CTCGAGTACAACATCCGGGTCTACTGCCTGCATGACACCCACGATGCACTCAAGGAGGTG 2043
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Db     958 CTCGAGTACAACATCCGGGTCTACTGCCTGCATGACACCCACGATGCACTCAAGGAGGTG 1017

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Db    1018 GTGCAGCTGGAGAAGCAGCTGGGGGGACAGCTGATCCAGGAGCCACGGGTCCTGCACTT- 1076

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Db    1077 AAGGACAGTTACCACAACCT--GCCCTATCATCCACGATGTGCCCAGCTCCCTGTGGAAG 1134

Qy    2164 AGTAAGCTCCTTGTCAGCTACCAGGAGATCCCCTTTTATCACATCTGGAATGGCACGCAG 2223
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Db    1135 AGTAAGCTCCTTGTCAGCTACCAGGAGATCCCCTTTTATCACATCTGGAATGGCACGCAG 1194

Qy    2224 CGGTACTTGCACTGCACCTTCACCCTGGAGCGTGTCAGCCCCAGCACTAGTGACCTGGCC 2283
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Db    1195 CGGTACTTGCACTGCACCTTCACCCTGGAGCGTGTCAGCCCCAGCACTAGTGACCTGGCC 1254

Qy    2284 TGCAAGCTGTGGGTGTGGCAGGTGGAGGGCGACGGGCAGAGCTTCAGCATCAACTTCAAC 2343
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Qy    2464 AGCCTGGACCCACCCTGTAGGCGGGGTGCCGACTGGCGGACTCTGGCCCAGAAACTCCAC 2523
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Db    1495 CTGGACAGCCATCTCAGCTTCTTTGCCTCCAAGCCCAGCCCCACAGCCATGATCCTCAAC 1554

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RESULT 10
US-10-256-702-2
; Sequence 2, Application US/10256702
; Publication No. US20030059859A1
;   GENERAL INFORMATION:
;   APPLICANT: Tessier-Lavigne, Marc
;   Leonardo, E. David

US-10-256-702-2

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    Query Match              57.6%;  Score 1552.4;  DB 14;  Length 1787;
    Best Local Similarity    98.5%;  Pred. No. 0;
    Matches 1651;  Conservative    0;  Mismatches   16;  Indels    9;  Gaps     8;

Qy      1025 GCAACTGTACCAGTGACCTCTGTGTACACAGTGCTTCTGGCCCTGAGGACGTGGCCCTCT 1084
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Db        60 ATGTGGGCCTCATCGCCGTGGCCGTCTGCCTGGTCCTGCTGCTGCTTGTCCTCATCCTCG 119

Qy      1145 TTTATTGCCGGAAGAAGGAGGGGCTGGACTCAGATGTGGCTGACTCGTCCATTCTCACCT 1204
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             ||||||||||||||||||||||| |||||||||||||||||||||||||||||||||||||
Db      180  CAGGCTTCCAGCCCGTCAGCATCTAAGCCCAGCAAAGCAGACAACCCCCATCTGCTCACC 239

Qy     1264  ATCCAGCCGGACCTCAGCACCACCACCACCACCTACCAGGGCAGTCTCTGTCCCCGGCAG 1323
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Db      240  ATCCAGCCGGACCTCAGCACCACCACCACCACCTACCAGGGCAGTCTCTGTCCCCGGCAG 299

Qy     1324  GATGGGCCCAGCCCCAAGTTCCAGCTCACCAATGGGCACCTGCTCAGCCCCCTGGGTGGC 1383
             ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db      300  GATGGGCCCAGCCCCAAGTTCCAGCTCACCAATGGGCACCTGCTCAGCCCCCTGGGTGGC 359

Qy     1384  GGCCGCCACACACTGCACCACAGCTCTCCCACCTCTGAGGCCGAGGAGTTCGTCTCCCGC 1443
             ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db      360  GGCCGCCACACACTGCACCACAGCTCTCCCACCTCTGAGGCCGAGGAGTTCGTCTCCCGC 419

Qy     1444  CTCTCCACCCAGAACTACTTCCGCTCCCTGCCCCGAGGCACCAGCAACATGACCTATGGG 1503
             ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db      420  CTCTCCACCCAGAACTACTTCCGCTCCCTGCCCCGAGGCACCAGCAACATGACCTATGGG 479

Qy     1504  ACCTTCAACTTCCTCGGGGGCCGGCTGATGATCCCTAATACAGGTATCAGCCTCCTCATC 1563
             ||||||||||||||||||||||||||||||||||||||||||| ||||||||||||||||
Db      480  ACCTTCAACTTCCTCGGGGGCCGGCTGATGATCCCTAATACAGGAATCAGCCTCCTCATC 539

Qy     1564  CCCCCAGATGCCATACCCCGAGGGAAGATCTATGAGATCTACCTCACGCTGCACAAGCCG 1623
             |||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db      540  CCCCCAGATGCCATACCCCGAGGGAAGATCTATGAGATCTACCTCACGCTGCACAAGCCG 599

Qy     1624  GAAGACGTGAGGTTGCCCCTAGCTGGCTGTCAGACCCTGCTGAGTCCCATCGTTAGCTGT 1683
             ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db      600  GAAGACGTGAGGTTGCCCCTAGCTGGCTGTCAGACCCTGCTGAGTCCCATCGTTAGCTGT 659

Qy     1684  GGACCCCCTGGCGTCCTGCTCACCCGGCCAGTCATCCTGGCTATGGACCACTGTGGGGAG 1743
             ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db      660  GGACCCCCTGGCGTCCTGCTCACCCGGCCAGTCATCCTGGCTATGGACCACTGTGGGGAG 719

Qy     1744  CCCAGCCCTGACAGCTGGAGCCTGCGCCTCAAAAAGCAGTCGTGCGAGGGCAGCTGGGAG 1803
             ||||||||||||||||||||||||    ||||||||||||||||||||||| |||||||||
Db      720  CCCAGCCCTGACAGCTGGAGCCTGGCCCTCAAAAAGCAGTCGTGCGAGGG-AGCTGGGAG 778

Qy     1804  GATGTGCTGCACCTGGGCGAGGAGGCGCCCTCCCACCTCTACTACTGCCAGCTGGAGGCC 1863
             |||||  ||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db      779  GATGT-CTGCACCTGGGCGAGGAGGCGCCCTCCCACCTCTACTACTGCCAGCTGGAGGCC 837

Qy     1864  AGTGCCTGCTACGTCTTCACCGAGCAGCTGGGCCGCTTTGCCCTGGTGGGAGAGGCCCTC 1923
             ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db      838  AGTGCCTGCTACGTCTTCACCGAGCAGCTGGGCCGCTTTGCCCTGGTGGGAGAGGCCCTC 897

Qy     1924  AGCGTGGCTGCCGCCAAGCGCCTCAAGCTGCTTCTGTTTGCGCCGGTGGCCTGCACCTCC 1983
             ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db      898  AGCGTGGCTGCCGCCAAGCGCCTCAAGCTGCTTCTGTTTGCGCCGGTGGCCTGCACCTCC 957

Qy     1984  CTCGAGTACAACATCCGGGTCTACTGCCTGCATGACACCCACGATGCACTCAAGGAGGTG 2043
             ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db      958  CTCGAGTACAACATCCGGGTCTACTGCCTGCATGACACCCACGATGCACTCAAGGAGGTG 1017
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Qy      2044 GTGCAGCTGGAGAAGCAGCTGGGGGGGACAGCTGATCCAGGAGCCACGGGTCCTGCACTTC 2103
             ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db      1018 GTGCAGCTGGAGAAGCAGCTGGGGGGGACAGCTGATCCAGGAGCCACGGGTCCTGCACTT- 1076

Qy      2104 AAGGACAGTTACCACAACCTGCGCCTATCCATCCACGATGTGCCCAGCTCCCTGTGGAAG 2163
             ||||||||||||||||||||  |||     |||||||||||||||||||||||||||||||
Db      1077 AAGGACAGTTACCACAACCT--GCCCTATCATCCACGATGTGCCCAGCTCCCTGTGGAAG 1134

Qy      2164 AGTAAGCTCCTTGTCAGCTACCAGGAGATCCCCTTTTATCACATCTGGAATGGCACGCAG 2223
             ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db      1135 AGTAAGCTCCTTGTCAGCTACCAGGAGATCCCCTTTTATCACATCTGGAATGGCACGCAG 1194

Qy      2224 CGGTACTTGCACTGCACCTTCACCCTGGAGCGTGTCAGCCCCAGCACTAGTGACCTGGCC 2283
             ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db      1195 CGGTACTTGCACTGCACCTTCACCCTGGAGCGTGTCAGCCCCAGCACTAGTGACCTGGCC 1254

Qy      2284 TGCAAGCTGTGGGTGTGGCAGGTGGAGGGCGACGGGCAGAGCTTCAGCATCAACTTCAAC 2343
             ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db      1255 TGCAAGCTGTGGGTGTGGCAGGTGGAGGGCGACGGGCAGAGCTTCAGCATCAACTTCAAC 1314

Qy      2344 ATCACCAAGGACACAAGGTTTGCTGAGCTGCTGGCTCTGGAGAGTGAAGCGGGGGTCCCA 2403
             ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db      1315 ATCACCAAGGACACAAGGTTTGCTGAGCTGCTGGCTCTGGAGAGTGAAGCGGGGGTCCCA 1374

Qy      2404 GCCCTGGTGGGCCCCAGTGCCTTCAAGATCCCCTTCCTCATTCGGCAGAAGATAATTTCC 2463
             |||||||||||||||||||||||||||||||||||||||||||||||||||||||||||,|
Db      1375 GCCCTGGTGGGCCCCAGTGCCTTCAAGATCCCCTTCCTCATTCGGCAGAAGATAATTTCC 1434

Qy      2464 AGCCTGGACCCACCCTGTAGGCGGGGTGCCGACTGGCGGACTCTGGCCCAGAAACTCCAC 2523
             ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db      1435 AGCCTGGACCCACCCTGTAGGCGGGGTGCCGACTGGCGGACTCTGGCCCAGAAACTCCAC 1494

Qy      2524 CTGGACAGCCATCTCAGCTTCTTTGCCTCCAAGCCCAGCCCCACAGCCATGATCCTCAAC 2583
             ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db      1495 CTGGACAGCCATCTCAGCTTCTTTGCCTCCAAGCCCAGCCCCACAGCCATGATCCTCAAC 1554

Qy      2584 CTGTGGGAGGCGCGGCACTTCCCCAACGGCAACCTCAGCCAGCTGGCTGCAGCAGTGGCT 2643
             ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db      1555 CTGTGGGAGGCGCGGCACTTCCCCAACGGCAACCTCAGCCAGCTGGCTGCAGCAGTGGCT 1614

Qy      2644 GGACTGGGCCAGCCAGACGCTGGCCTC-TTCACAGTG-TCGGAGGCTGAGTGCTGA 2697
             ||        |||||||   ||||  ||||  ||  ||||||||||| ||||||||||||||||||||
Db      1615 GGGACTGGCCAGCAGGACGGTGGCTTCTTTCACAGTGTTCGGAGGCTGAGTGCTGA 1670
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RESULT 11
US-10-296-115-365

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   Query Match          44.5%;  Score 1200.6;  DB 17;  Length 1321;
   Best Local Similarity  98.0%;  Pred. No. 8.1e-301;
   Matches 1289;  Conservative    0;  Mismatches  19;  Indels   7;  Gaps    7;

Qy     1390 CACACACTGCACCACAGCTCTCCCACCTCTGAGGCCGAGGAGTTCGTCTCCCGCCTCTCC 1449
            ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        1 CACACACTGCACCACAGCTCTCCCACCTCTGAGGCCGAGGAGTTCGTCTCCCGCCTCTCC 60

Qy     1450 ACCCAGAACTACTTCCGCTCCCTGCCCCGAGGCACCAGCAACATGACCTATGGGACCTTC 1509
            ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db       61 ACCCAGAACTACTTCCGCTCCCTGCCCCGAGGCACCAGCAACATGACCTATGGGACCTTC 120

Qy     1510 AACTTCCTCGGGGGCCGGCTGATGATCCCTAATACAGGTATCAGCCTCCTCATCCCCCCA 1569
            |||||||||||||||||||||||||||||||||||||||| |||||||||||||||||||
Db      121 AACTTCCTCGGGGGCCGGCTGATGATCCCTAATACAGGAATCAGCCTCCTCATCCCCCCA 180

Qy     1570 GATGCCATACCCCGAGGGAAGATCTATGAGATCTACCTCACGCTGCACAAGCCGGAAGAC 1629
            ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db      181 GATGCCATACCCCGAGGGAAGATCTATGAGATCTACCTCACGCTGCACAAGCCGGAAGAC 240

Qy     1630 GTGAGGTTGCCCCTAGCTGGCTGTCAGACCCTGCTGAGTCCCATCGTTAGCTGTGGACCC 1689
            ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db      241 GTGAGGTTGCCCCTAGCTGGCTGTCAGACCCTGCTGAGTCCCATCGTTAGCTGTGGACCC 300

Qy     1690 CCT-GGCGTCCTGCTCACCCGGCCAGTCATCCT-GGCTATGGACCACTGT-GGGGAGCCC 1746
            ||| |||||||||| ||||||||||||||||| || ||||||||||| .|||||||||
Db      301 CCTGGGCGTCCTGCTTACCCGGCCAGTCATCCTGGGGTATGGACCACTGTGGGGGAGCCC 360

Qy     1747 AGCCCTGACAGCT-GGAGCCTGCGCCTCAAAAAGCAGTCGTGCGAGGGCAGCTGGGAGGA 1805
            ||||||||||||| ||||||||||||||||||||||||||||||||||||||||||||||
Db      361 AGCCCTGACAGCTGGGAGCCTGCGCCTCAAAAAGCAGTCGTGCGAGGGCAGCTGGGAGGA 420

Qy     1806 TGTGCTGCACCTGGGCGAGGAGGCGCCCTCCCACCTCTACTACTGCCAGCTGGAGGCCAG 1865
            ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db      421 TGTGCTGCACCTGGGCGAGGAGGCGCCCTCCCACCTCTACTACTGCCAGCTGGAGGCCAG 480

Qy     1866 TGCCTGCTACGTCTTCACCGAGCAGCTGGGCCGCTTTGCCCTGGTGGGAGAGGCCCTCAG 1925
            ||||||||||||||||||||||||||| ||||||| ||||||||||||||||||||||||
Db      481 TGCCTGCTACGTCTTCACCGAGCAGCTGAGCCGCTATGCCCTGGTGGGAGAGGCCCTCAG 540

Qy     1926 CGTGGCTGCCGCCAAGCGCCTCAAGCTGCTTCTGTTTGCGCCGGTGGCCTGCACCTCCCT 1985
            ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db      541 CGTGGCTGCCGCCAAGCGCCTCAAGCTGCTTCTGTTTGCGCCGGTGGCCTGCACCTCCCT 600

Qy     1986 CGAGTACAACATCCGGGTCTACTGCCTGCATGACACCCACGATGCACTCAAGGAGGTGGT 2045
            |||||||||| | ||||||||||||||||||| ||||||||||| | |||||
Db      601 CGAGTACAACATACTGGTCTACTGCCTGCATGACACTCACGATGCACTCAACGTAGTGGT 660
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Qy    2046  GCAGCTGGAGAAGCAGCTGGGGGGGACAGCTGATCCAGGAGCCACGGGTCCTGCACTTCAA 2105
            ||||||||||||||||||   ||||||||||||||||||||||| |||  ||||||||||
Db     661  GCAGCTGGAGAAGCAGCTGCAGGGACAGCTGATCCAGGAGCCACTGGTACTGCACTTCAA 720

Qy    2106  GGACAGTTACCACAACCTGCGCCTATCCATCCACGATGTGCCCAGCTCCCTGTGGAAGAG 2165
            ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db     721  GGACAGTTACCACAACCTGCGCCTATCCATCCACGATGTGCCCAGCTCCCTGTGGAAGAG 780

Qy    2166  TAAGCTCCTTGTCAGCTACCAGGAGATCCCCTTTTATCACATCTGGAATGGCACGCAGCG 2225
            ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db     781  TAAGCTCCTTGTCAGCTACCAGGAGATCCCCTTTTATCACATCTGGAATGGCACGCAGCG 840

Qy    2226  GTACTTGCACTGCACCTTCACCCTGGAGCGTGTCAGCCCCAGCACTAGTGACCTGGCCTG 2285
            ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db     841  GTACTTGCACTGCACCTTCACCCTGGAGCGTGTCAGCCCCAGCACTAGTGACCTGGCCTG 900

Qy    2286  CAAGCTGTGGGTGTGGCAGGTGGAGGGCGACGGGCAGAGCTTCAGCATCAACTTCAACAT 2345
            ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db     901  CAAGCTGTGGGTGTGGCAGGTGGAGGGCGACGGGCAGAGCTTCAGCATCAACTTCAACAT 960

Qy    2346  CACCAAGGACACAAGGTTTGCTGAGCTGCTGGCTCTGGAGAGTGAAGCGGGGGTCCCAGC 2405
            ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db     961  CACCAAGGACACAAGGTTTGCTGAGCTGCTGGCTCTGGAGAGTGAAGCGGGGGTCCCAGC 1020

Qy    2406  CCTGGTGGGCCCCAGTGCCTTCAAGATCCCCTTCCTCATTCGGCAGAAGATAATTTCCAG 2465
            ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    1021  CCTGGTGGGCCCCAGTGCCTTCAAGATCCCCTTCCTCATTCGGCAGAAGATAATTTCCAG 1080

Qy    2466  CCTGGACCCACCCTGTAGGCGGGGTGCCGACTGGCGGACTCTGGCCCAGAAACTCCACCT 2525
            ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    1081  CCTGGACCCACCCTGTAGGCGGGGTGCCGACTGGCGGACTCTGGCCCAGAAACTCCACCT 1140

Qy    2526  GGACAGCCATCTCAGCTTCTTTGCCTCCAAGCCCAGCCCCACAGCCATGATCCTCAACCT 2585
            ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    1141  GGACAGCCATCTCAGCTTCTTTGCCTCCAAGCCCAGCCCCACAGCCATGATCCTCAACCT 1200

Qy    2586  GTGGGAGGCGCGGCACTTCCCCAACGGCAACCTCAGCCAGCTGGCTGCAGCAGTGGCT-G 2644
            |||||||||||||||||||||||||||||||||||||||||||||||||||||||||| |
Db    1201  GTGGGAGGCGCGGCACTTCCCCAACGGCAACCTCAGCCAGCTGGCTGCAGCAGTGGCTGG 1260

Qy    2645  GACTGGGCCAGCCAGACGCTGGCCTC-TTCACAGTG-TCGGAGGCTGAGTGCTGA 2697
            ||||||||||| |||| |||| || |||||||| ||||||||||||||||||
Db    1261  GACTGGGCCAGCAGGACGGTGGCTTCTTTCACAGTGTTCGGAGGCTGAGTGCTGA 1315
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RESULT 12
US-10-087-684-1
; Sequence 1, Application US/10087684
; Publication No. US20040029116A1
; GENERAL INFORMATION:
; APPLICANT: Edinger, Shlomit R.
; APPLICANT: MacDougall, John R.
; APPLICANT: Millet, Isabelle
; APPLICANT: Ellerman, Karen
; APPLICANT: Stone, David J.
; APPLICANT: Grosse, William M.

```
  Query Match              34.7%;  Score 936.2;  DB 17;  Length 2860;
  Best Local Similarity    61.7%;  Pred. No. 2.8e-232;
  Matches 1662;  Conservative    0;  Mismatches 938;  Indels   93;  Gaps      7;

Qy        98 ACCCAGTGCCTGGTGCCAACCCGGACCTGCTTCCCCACTTCCTGGTGGAGCCCGAGGATG 157
             || |  | ||     ||     | || | ||| ||| |||||||| |||||| |||| |
Db       168 ACTCCTTCCCGTCAGCGCCAGCAGAGCCGCTGCCCTACTTCCTGCAGGAGCCACAGGACG 227

Qy       158 TGTACATCGTCAAGAACAAGCCAGTGCTGCTTGTGTGCAAGGCCGTGCCCGCCACGCAGA 217
             ||||| || |||||||||||| ||| |||     ||| ||| | |||||||||| ||||
Db       228 CCTACATTGTGAAGAACAAGCCTGTGGAGCTCCGCTGCCGCGCCTTCCCCGCCACACAGA 287
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Qy    218 TCTTCTTCAAGTGCAACGGGGAGTGGGTGCGCCAGGTGGACCACGTGATCGAGCGCAGCA 277
          ||| ||||||||||||||| |||||||| ||||| ||||||.||| |  ||    ||
Db    288 TCTACTTCAAGTGCAACGGCGAGTGGGTCAGCCAGAACGACCACGTCACACAGGAAGGCC 347

Qy    278 CAGACGGGAGCAGTGGGCTGCCCACCATGGAGGTCCGCATTAATGTCTCAAGGCAGCAGG 337
           || | | || || |||| ||||| | || | || |||||||||
Db    348 TGGATGAGGCCACCGGTCTGCGGGTGCGCGAGGTGCAGATCGAGGTGTCGCGGCAGCAGG 407

Qy    338 TCGAGAAGGTGTTCGGGCTGGAGGAATACTGGTGCCAGTGCGTGGCATGGAGCTCCTCGG 397.
          | ||| || | || ||||||||||| |||||||||||||||||| ||||||||||| |||
Db    408 TGGAGGAGCTCTTTGGGCTGGAGGATTACTGGTGCCAGTGCGTGGCCTGGAGCTCCGCGG 467

Qy    398 GCACCACCAAGAGTCAGAAGGCCTACATCCGCATAGCCAGATTGCGCAAGAACTTCGAGC 457
          ||||||||||||||      |||||| ||||||| |||      ||||||||||||||| |
Db    468 GCACCACCAAGAGTCGCCGAGCCTACGTCCGCATCGCCTACCTGCGCAAGAACTTCGATC 527

Qy    458 AGGAGCCGCTGGCCAAGGAGGTGTCCCTGGAGCAGGGCATCGTGCTGCCCTGCCGTCCAC 517
          ||||||| |||| |||||||||| ||||||| ||||||| || |    | | |||| |||| || |
Db    528 AGGAGCCTCTGGGCAAGGAGGTGCCCCTGGACCATGAGGTTCTCCTGCAGTGCCGCCCGC 587

Qy    518 CGGAGGGCATCCCTCCAGCCGAGGTGGAGTGGCTCCGGAACGAGGACCTGGTGGACCCGT 577
          |||||||. | ||| |||||||||||| |||||| ||| ||||| | | |||||
Db    588 CGGAGGGGGTGCCTGTGGCCGAGGTGGAATGGCTCAAGAATGAGGATGTCATCGACCCCA 647

Qy    578 CCCTGGACCCCAATGTATACATCACGCGGGAGCACAGCCTGGTGGTGCGACAGGCCCGCC 637
          ||| |||| |||| |       |||     || ||||| |||  |  | || |||||||||||.
Db    648 CCCAGGACACCAACTTCCTGCTCACCATCGACCACAACCTCATCATCCGCCAGGCCCGCC 707

Qy    638 TTGCTGACACGGCCAACTACACCTGCGTGGCCAAGAACATCGTGGCACGTCGCCGCAGCG 697
          |  |  ||||| ||||||||| |||||||||||||||||||||||||||||||     ||||| |||
Db    708 TGTCGGACACTGCCAACTATACCTGCGTGGCCAAGAACATCGTGGCCAAACGCCGGAGCA 767

Qy    698 CCTCCGCTGCTGTCATCGTCTACGTGAACGGTGGGTGGTCGACGTGGACCGAGTGGTCCG 757
          || |  || | ||||||||||||||||| || || |||||| | ||| | ||||||||
Db    768 CCACTGCCACCGTCATCGTCTACGTGAATGGCGGCTGGTCCAGCTGGGCAGAGTGGTCAC 827

Qy    758 TCTGCAGCGCCAGCTGTGGGCGCGGCTGGCAGAAACGGAGCCGGAGCTGCACCAACCCGG 817
           ||||   |  | ||||||| ||  |||||||||| || |  ||||| |||||||||||| |
Db    828 CCTGCTCCAACCGCTGTGGCCGAGGCTGGCAGAAGCGCACCCGGACCTGCACCAACCCCG 887

Qy    818 CGCCTCTCAACGGGGGCGCTTTCTGTGAGGGGCAGAATGTCCAGAAAACAGCCTGCGCCA 877
          | || ||||||||| || || ||||| ||||| |||      ||||||| || ||||||  |||
Db    888 CTCCACTCAACGGAGGGGCCTTCTGCGAGGGCCAGGCATTCCAGAAGACCGCCTGCACCA 947

Qy    878 CCCTGTGCCCAGTAGACGGCAGCTGGAGCCCGTGGAGCAAGTGGTCGGCCTGTGGGCTGG 937
          || | ||||||||| || || |||| ||||||||||||||||| |||||  |     |
Db    948 CCATCTGCCCAGTCGATGGGGCGTGGACGGAGTGGAGCAAGTGGTCAGCCTGCAGCACTG 1007

Qy    938 ACTGCACCCACTGGCGGAGCCGTGAGTGCTCTGACCCAGCACCCCGCAACGGAGGGGAGG 997
          | ||  ||||||||||| |||||| ||||| |||||  |  || |||||| ||||||||  |
Db   1008 AGTGTGCCCACTGGCGTAGCCGCGAGTGCATGGCGCCCCCACCCCAGAACGGAGGCCGTG 1067

Qy    998 AGTGCCAGGGCACTGACCTGGACACCCGCAACTGTACCAGTGACCTCTGTGTACACAGTG 1057
          | ||| ||  || || || || |  ||  |||||  ||||| || || || | ||  |
Db   1068 ACTGCAGCGGGACGCTGCTCGACTCTAAGAACTGCACAGATGGGCTGTGCATGCAACTGG 1127

Qy   1058 CTTCTGGCCCTGAGGACGTGGCCCTCTATGTGGGCCTCATCGCCGTGGCCGTCTGCCTGG 1117
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                        |       || |   | |           | |   | |   | ||| |   |||   | |||
Db    .1128 AGGCCTCAGGGGATGCGGCGCTGTATGCGGGGCTCGTGGTGGCCATCTTCGTGGTCGTGG 1187

Qy     1118 TCCTGCTGCTGCTTGTCCTCATCCTCGTTTATTGCCGGAAGAAGGAGGGGCTGGACTCAG 1177
              | |||  || ||   | ||    |  | | || ||  |||| ||      |   | ||| |||
Db     1188 CAATCCTCATGGCGGTGGGGGTGGTGGTGTACCGCCGCAACTGCCGTGACTTCGACACAG 1247

Qy     1178 ATGTGGCTGACTCGTCCATT---CTCACCTCAGGCTTCCAGCCCGTCAGCATCAAGCCCA 1234
              | |  ||||||| ||   |    || ||    || |||||| |||||| | | ||| |
Db     1248 ACATCACTGACTCATCTGCTGCCCTGACTGGTGGTTTCCACCCCGTCAACTTTAAGACGG 1307

Qy     1235 GCAAAGCAGACAACCCCCATCTGCT------CACCATCCAGCCGGACCTCAGCACCACCA 1288
              | |    |||| || || ||    | | |    || ||||| |   ||| |
Db     1308 CAAGGCCCAGTAACCCGCAGCTCCTACACCCCTCTGTGCCTCCTGACCTGACAGCCAGCG 1367

Qy     1289 CCACCACCTACCAGGGCAGTCTCTGTCCCCGGCAGGA---------------------- 1325
              ||   || |||||  ||     | | | ||| ||||||
Db     1368 CCGGCATCTACCGCGGACCCGTGTATGCCCTGCAGGACTCCACCGACAAAATCCCCATGA 1427

Qy     1326 ---------------------TGGGCCCAGCCCCAAGTTCCAGCTCACCA--------- 1354
                                   ||||||| ||| || || || ||   ||
Db     1428 CCAACTCTCCTCTGCTGGACCCCCTTACCCAGCCTTAAGGTCAAGGTCTACAGCTCCAGCA 1487

Qy     1355 --ATGGGCACCTGCTCAGCCCCCTGGGTGGCGGCCGCCACAC----------ACTGCACC 1402
                | |||| |   | ||| ||     | | ||| | |    |         | ||
Db     1488 CCACGGGCTCTGGGCCAGGCCTGGCAGATGGGGCTGACCTGCTGGGGGTCTTGCCGCCTG 1547

Qy     1403 ACAGCTCTCCCACCTCTGAGGCCGAGGAGTTCGTCTCCCGCCTCTCCACCCAGAACTACT 1462
              || | || | | | |   ||| ||| |  | || ||   |    | |
Db     1548 GCACATACCCTAGCGATTTCGCCCGGGACACCCACTTCCTGCACCTGCGCAGCGCCAGCC 1607

Qy     1463 TCCGCTC---------------CCTGCCCCGAGGCACCAGCAACATGACCTATGGGACCT 1507
              || | ||              |||||||||||| | |   | | ||     |   || ||||
Db     1608 TCGGTTCCCAGCAGCTCTTGGGCCTGCCCCGAGACCCAGGGAGCAGCGTCAGCGGCACCT 1667

Qy     1508 TCAACTTCCTCGGGGGCCGGCTGATGATCCCTAATACAGGTATCAGCCTCCTCATCCCCC 1567
              |   || ||| || || ||   |||| |   ||||||     |||||  |||||| | ||  | |||
Db     1668 TTGGCTGCCTGGGTGGGAGGCTCAGCATCCCCGGCACAGGGGTCAGCTTGCTGGTGCCCA 1727

Qy     1568 CAGATGCCATACCCCGAGGGAAGATCTATGAGATCTACCTCACGCTGCACAAGCCGGAAG 1627
              |  ||||| ||||   || ||| |||| ||||| || ||      |  ||||| | |||
Db     1728 ATGGAGCCATTCCCCAGGGCAAGTTCTACGAGATGTATCTACTCATCAACAAGGCAGAAA 1787

Qy     1628 ACGTGAGGTTGCCCCTAGCTGGCTGTCAGACCCTGCTGAGTCCCATCGTTAGCTGTGGAC 1687
              | |  ||       || ||   ||     |||||  |  |||| |||   || | |||||||||
Db     1788 GTACCCTGCCGCTTTCAGAAGGGACCCAGACAGTATTGAGCCCCTCGGTGACCTGTGGAC 1847

Qy     1688 CCCCTGGCGTCCTGCTCACCCGGCCAGTCATCCTGGCTATGGACCACTGTGGGGAGCCCA 1747
              || |  |||  |||||||    ||| ||  ||||||||| |  ||| |||||||||  ||   ||
Db     1848 CCACAGGCCTCCTGCTGTGCCGCCCCGTCATCCTCACCATGCCCCACTGTGCCGAAGTCA 1907

Qy     1748 GCCCTGACAGCTGGAGCCTGCGCCTCAAAAAGCAGTCGTGCGAGGGCAGCTGGGAGGATG 1807
              | |    |||||| | |   ||||| |  ||| |    | ||||| || ||||||||| |
Db     1908 GTGCCCGTGACTGGATCTTTCAGCTCAAGACCCAGGCCCACCAGGGCCACTGGGAGGAGG 1967

Qy     1808 TGCTGCACCTGGGCGAGGAGGCGCCCTCCCACCTCTACTACTGCCAGCTGGAGGCCAGTG 1867
              || ||  ||||||  |||||| ||||||| | |        |  | || |||||||||||||||||||||| |||| |
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Db   1968 TGGTGACCCTGGATGAGGAGACCCTGAACACACCCTGCTACTGCCAGCTGGAGCCCAGGG 2027

Qy   1868 CCTGCTACGTCTTCACCGAGCAGCTGGGCCGCTTTGCCCTGGTGGGAGAGGCCCTCAGCG 1927
          ||||  || || |    || ||||||||||  || |    |   ||| ||| ||      |
Db   2028 CCTGTCACATCCTGCTGGACCAGCTGGGCACCTACGTGTTCACGGGCGAGTCCTATTCCC 2087

Qy   1928 TGGCTGCCGCCAAGCGCCTCAAGCTGCTTCTGTTTGCGCCGGTGGCCTGCACCTCCCTCG 1987
           | || | ||||||| ||| ||||||    | || || || |    |||||||||||| |
Db   2088 GCTCAGCAGTCAAGCGGCTCCAGCTGGCCGTCTTCGCCCCCGCCCTCTGCACCTCCCTGG 2147

Qy   1988 AGTACAACATCCGGGTCTACTGCCTGCATGACACCCACGATGCACTCAAGGAGGTGGTGC 2047
          |||||| | |||||||||||||||||||||| | |||||| |  |  ||||| |||||||||| ||
Db   2148 AGTACAGCCTCCGGGTCTACTGCCTGGAGGACACGCCTGTAGCACTGAAGGAGGTGCTGG 2207

Qy   2048 AGCTGGAGAAGCAGCTGGGGGGACAGCTGATCCAGGAGCCACGGGTCCTGCACTTCAAGG 2107
          ||||||||  |     ||||| ||| | |  | || | |||||||         ||   ||||||||
Db   2208 AGCTGGAGCGGACTCTGGGCGGATACTTGGTGGAGGAGCCGAAACCGCTAATGTTCAAGG 2267

Qy   2108 ACAGTTACCACAACCTGCGCCTATCCATCCACGATGTGCCCAGCTCCCTGTGGAAGAGTA 2167
          |||||||||||||||||||||||  ||| |||| ||  |  | ||    ||| |||| ||| |
Db   2268 ACAGTTACCACAACCTGCGCCTCTCCCTCCATGACCTCCCCCATGCCCATTGGAGGAGCA 2327

Qy   2168 AGCTCCTTGTCAGCTACCAGGAGATCCCCTTTTATCACATCTGGAATGGCACGCAGCGGT 2227
          ||||  || |  ||  ||||||||||||||||| |||||||| |||| ||||| |||  | |
Db   2328 AGCTGCTGGCCAAATACCAGGAGATCCCCTTCTATCACATTTGGAGTGGCAGCCAGAAGG 2387

Qy   2228 ACTTGCACTGCACCTTCACCCTGGAGCGTGTCAGCCCCAGCACTAGTGACCTGGCCTGCA 2287
           | | ||||||||| |||||||||||||| |    |||     | | | || || ||||||
Db   2388 CCCTCCACTGCACTTTCACCCTGGAGAGGCACAGCTTGGCCTCCACAGAGCTCACCTGCA 2447

Qy   2288 AGCTGTGGGTGTGGCAGGTGGAGGGCGACGGGCAGAGCTTCAGCATCAACTTCAACATCA 2347
          || |  || ||| |||| ||||| || || || |||| |||   |  | |    || |
Db   2448 AGATCTGCGTGCGGCAAGTGGAAGGGGAGGGCCAGATATTCCAGCTGCATACCACTCTGG 2507

Qy   2348 CCAAG---GACACAAGGTTTGCTGAGCTGCTGGCTCTGGAGAGTGAAGCGGGGGTCCCAG 2404
          | | ||      | | |     ||  ||    | |       || |          ||| |
Db   2508 CAGAGACACCTGCTGGCTCCCTGGACACTCTCTGCTCTGCCCCTGGCAGCACTGTCACCA 2567

Qy   2405 CCCTGGTGGGCCCCAGTGCCTTCAAGATCCCCTTCCTCATTCGGCAGAAGATAATTTCCA 2464
          ||| | |||| ||    ||||||||||||| |    ||| || ||||||||||        ||
Db   2568 CCCAGCTGGGACCTTATGCCTTCAAGATCCCACTGTCCATCCGCCAGAAGATATGCAACA 2627

Qy   2465 GCCTGGACCCACCCTGTAGGCGGGGTGCCGACTGGCGGACTCTGGCCCAGAAACTCCACC 2524
          ||||  ||  |  |||     |||||  |||||||||||  | || |||||| |||  |||
Db   2628 GCCTAGATGCCCCCAACTCACGGGGCAATGACTGGCGGATGTTAGCACAGAAGCTCTCTA 2687

Qy   2525 TGGACAGCCATCTCAGCTTCTTTGCCTCCAAGCCCAGCCCCACAGCCATGATCCTCAACC 2584
          |||||  |    | || |   | ||||||||| |||| | |||||||| |    ||||||| |||
Db   2688 TGGACCGGTACCTGAATTACTTTGCCACCAAAGCGAGCCCCACGGGTGTGATCCTGGACC 2747

Qy   2585 TGTGGGAGGCGCGGCACTTCCCCAACGGCAACCTCAGCCAGCTGGCTGCAGCAGTGGCTG 2644
          | ||||| || | |||    | || || |  ||||||  |||||        || ||| |
Db   2748 TCTGGGAAGCTCTGCAGCAGGACGATGGGGACCTCAACAGCCTGGCGAGTGCCTTGGAGG 2807

Qy   2645 GACTGGGCCAGCCAGACGCTGGCCTCTTCACAGTGTCGGAGGCTGAGTGCTGA 2697
          |||||| ||   ||      |         |    | ||| | |||||||
Db   2808 AGATGGGCAAGAGTGAGATGCTGGTGGCTGTGGCCACCGACGGGGACTGCTGA 2860
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; Sequence 3, Application US/10087684
; Publication No. US20040029116A1
; GENERAL INFORMATION:
; APPLICANT: Edinger, Shlomit R.
; APPLICANT: MacDougall, John R.
; APPLICANT: Millet, Isabelle
; APPLICANT: Ellerman, Karen
; APPLICANT: Stone, David J.
; APPLICANT: Grosse, William M.
; APPLICANT: Lepley, Denise M.
; APPLICANT: Rieger, Daniel K.
; APPLICANT: Burgess, Cat250 Cathereine E.
; APPLICANT: Casman, Stacie, J.
; APPLICANT: Spytek, Kimberly A.
; APPLICANT: Boldog, Ferenc L.
; APPLICANT: Li, Li
; APPLICANT: Padigaru, Muralidhara
; APPLICANT: Mishra, Vishnu
; APPLICANT: Shenoy, Suresh G.
; APPLICANT: Rastelli, Luca
; APPLICANT: Tchernev, Velizar T.
; APPLICANT: Vernet, Corine A.M.
; APPLICANT: Zerhusen, Bryan D.
; APPLICANT: Malyankar, Uriel M.
; APPLICANT: Guo, Xiaojia
; APPLICANT: Miller, Charles E.
; APPLICANT: Gangolli, Esha A.
; TITLE OF INVENTION: PROTEINS AND NUCLEIC ACIDS ENCODING SAME
; FILE REFERENCE: 21402-214 CIP
; CURRENT APPLICATION NUMBER: US/10/087,684
; CURRENT FILING DATE: 2003-03-10
; PRIOR APPLICATION NUMBER: 60/253,834
; PRIOR FILING DATE: 2000-11-29
; PRIOR APPLICATION NUMBER: 60/250,926
; PRIOR FILING DATE: 2000-11-30
; PRIOR APPLICATION NUMBER: 60/264,180
; PRIOR FILING DATE: 2001-01-25
; PRIOR APPLICATION NUMBER: 60/274,194
; PRIOR FILING DATE: 2001-03-08
; PRIOR APPLICATION NUMBER: 60/313,656
; PRIOR FILING DATE: 2001-08-20
; PRIOR APPLICATION NUMBER: 60/327,456
; PRIOR FILING DATE: 2001-10-05
; NUMBER OF SEQ ID NOS: 220
; SOFTWARE: CuraSeqList version 0.1
; SEQ ID NO 3
;   LENGTH: 2860
;   TYPE: DNA
;   ORGANISM: Homo sapiens
;   FEATURE:
; ' NAME/KEY: CDS
;   LOCATION: (59)..(2857)

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Query Match            34.7%;  Score 936.2;  DB 17;  Length 2860;
Best Local Similarity  61.7%;  Pred. No. 2.8e-232;
Matches 1662;  Conservative     0;  Mismatches  938;  Indels   93;  Gaps      7;


Qy       98 ACCCAGTGCCTGGTGCCAACCCGGACCTGCTTCCCCACTTCCTGGTGGAGCCCGAGGATG 157
            || |  | ||    ||        | || | ||| ||| |||||||| |||||| |||| |
Db      168 ACTCCTTCCCGTCAGCGCCAGCAGAGCCGCTGCCCTACTTCCTGCAGGAGCCACAGGACG 227


Qy      158 TGTACATCGTCAAGAACAAGCCAGTGCTGCTTGTGTGCAAGGCCGTGCCCGCCACGCAGA 217
            ||||| || ||||||||||||| ||| |||| ||| ||| |||||||||| ||||
Db      228 CCTACATTGTGAAGAACAAGCCTGTGGAGCTTCGCTGCCGCGCCTTCCCCGCCACACAGA 287


Qy      218 TCTTCTTCAAGTGCAACGGGGAGTGGGTGCGCCAGGTGGACCACGTGATCGAGCGCAGCA 277
            ||| |||||||||||||||| |||||||| ||||| |||||||| |  ||      ||
Db      288 TCTACTTCAAGTGCAACGGCGAGTGGGTCAGCCAGAACGACCACGTCACACAGGAAGGCC 347


Qy      278 CAGACGGGAGCAGTGGGCTGCCCACCATGGAGGTCCGCATTAATGTCTCAAGGCAGCAGG 337
            || | |  || || |||      ||||| | || | || || |||||||||
Db      348 TGGATGAGGCCACCGGCCTGCGGGTGCGCGAGGTGCAGATCGAGGTGTCGCGGCAGCAGG 407


Qy      338 TCGAGAAGGTGTTCGGGCTGGAGGAATACTGGTGCCAGTGCGTGGCATGGAGCTCCTCGG 397
            | ||| || | || |||||||||||| ||||||||||||||||||||| ||||||||| | |
Db      408 TGGAGGAGCTCTTTGGGCTGGAGGATTACTGGTGCCAGTGCGTGGCCTGGAGCTCCGCAG 467


Qy      398 GCACCACCAAGAGTCAGAAGGCCTACATCCGCATAGCCAGATTGCGCAAGAACTTCGAGC 457
            ||||||||||||||    |||||| ||||||| |||      |||||||||||||||| |
Db      468 GCACCACCAAGAGTCGCCGAGCCTACGTCCGCATCGCCTACCTGCGCAAGAACTTCGATC 527


Qy      458 AGGAGCCGCTGGCCAAGGAGGTGTCCCTGGAGCAGGGCATCGTGCTGCCCTGCCGTCCAC 517
            ||||||| ||||  ||||||||| |||||||||| ||||| || |    |  | |||| || |
Db      528 AGGAGCCTCTGGGCAAGGAGGTGCCCCTGGACCATGAGGTTCTCCTGCAGTGCCGCCCGC 587


Qy      518 CGGAGGGCATCCCTCCAGCCGAGGTGGAGTGGCTCCGGAACGAGGACCTGGTGGACCCGT 577
            |||||||  |  ||| |||||||||||| ||||||| ||| ||||| |  | |||||
Db      588 CGGAGGGGGGTGCCTGTGGCCGAGGTGGAATGGCTCAAGAATGAGGATGTCATCGACCCCA 647


Qy      578 CCCTGGACCCCAATGTATACATCACGCGGGAGCACAGCCTGGTGGTGCGACAGGCCCGCC 637
            ||| |||| |||| |    ||||    || |||| ||| | | || |||||||||||
Db      648 CCCAGGACACCAACTTCCTGCTCACCATCGACCACAACCTCATCATCCGCCAGGCCCGCC 707


Qy      638 TTGCTGACACGGCCAACTACACCTGCGTGGCCAAGAACATCGTGGCACGTCGCCGCAGCG 697
            |  |  |||||| |||||||| ||||||||||||||||||||||||||||    ||||| |||
Db      708 TGTCGGACACTGCCAACTATACCTGCGTGGCCAAGAACATCGTGGCCAAACGCCGGAGCA 767


Qy      698 CCTCCGCTGCTGTCATCGTCTACGTGAACGGTGGGTGGTCGACGTGGACCGAGTGGTCCG 757
            || |  || | ||||||||||||||||||| || || |||||| |   ||| | ||||||||
Db      768 CCACTGCCACCGTCATCGTCTACGTGAATGGCGGCTGGTCCAGCTGGGCAGAGTGGTCAC 827


Qy      758 TCTGCAGCGCCAGCTGTGGGCGCGGCTGGCAGAAACGGAGCCGGAGCTGCACCAACCCGG 817
            ||||  |  | |||||||| || |||||||||| |  |||||| ||||||||||||||| |
Db      828 CCTGCTCCAACCGCTGTGGCCGAGGCTGGCAGAAGCGCACCCGGACCTGCACCAACCCCG 887


Qy      818 CGCCTCTCAACGGGGGCGCTTTCTGTGAGGGGCAGAATGTCCAGAAAACAGCCTGCGCCA 877
            | || |||||||| || || ||||| ||||| ||||| |||       ||||||| || |||||| |||
Db      888 CTCCACTCAACGGAGGGGCCTTCTGCGAGGGCCAGGCATTCCAGAAGACCGCCTGCACCA 947
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Qy    878 CCCTGTGCCCAGTAGACGGCAGCTGGAGCCCGTGGAGCAAGTGGTCGGCCTGTGGGCTGG 937
          || | |||||||| || ||    ||||      ||||||||||||||| ||||| |    |
Db    948 CCATCTGCCCAGTCGATGGGGCGTGGACGGAGTGGAGCAAGTGGTCAGCCTGCAGCACTG 1007

Qy    938 ACTGCACCCACTGGCGGAGCCGTGAGTGCTCTGACCCAGCACCCCGCAACGGAGGGGAGG 997
          | || ||||||||| ||||| |||||| | ||  |||||| ||||||| |
Db   1008 AGTGTGCCCACTGGCGTAGCCGCGAGTGCATGGCGCCCCCACCCCAGAACGGAGGCCGTG 1067

Qy    998 AGTGCCAGGGCACTGACCTGGACACCCGCAACTGTACCAGTGACCTCTGTGTACACAGTG 1057
          | ||| || || || ||| | |||| |     ||||| || || || || | ||    |
Db   1068 ACTGCAGCGGGACGCTGCTCGACTCTAAGAACTGCACAGATGGGCTGTGCATGCAACTGG 1127

Qy   1058 CTTCTGGCCCTGAGGACGTGGCCCTCTATGTGGGCCTCATCGCCGTGGCCGTCTGCCTGG 1117
          |        || | | |         |  | |  | | | |  | ||| |   |  | |||
Db   1128 AGGCCTCAGGGGATGCGGCGCTGTATGCGGGGCTCGTGGTGGCCATCTTCGTGGTCGTGG 1187

Qy   1118 TCCTGCTGCTGCTTGTCCTCATCCTCGTTTATTGCCGGAAGAAGGAGGGGCTGGACTCAG 1177
          | || || ||   |    | |  | |||| || ||| ||   |    | ||| ||| |||
Db   1188 CAATCCTCATGGCGGTGGGGGGTGGTGGTGTACCGCCGCAACTGCCGTGACTTCGACACAG 1247

Qy   1178 ATGTGGCTGACTCGTCCATT---CTCACCTCAGGCTTCCAGCCCGTCAGCATCAAGCCCA 1234
          | | ||||||| ||   |   || || ||||| ||||||| |  | ||| |
Db   1248 ACATCACTGACTCATCTGCTGCCCTGACTGGTGGTTTCCACCCCGTCAACTTTAAGACGG 1307

Qy   1235 GCAAAGCAGACAACCCCCATCTGCT------CACCATCCAGCCGGACCTCAGCACCACCA 1288
          | | |     ||||| || || ||     | | | || ||||| |   ||| |
Db   1308 CAAGGCCCAGTAACCCGCAGCTCCTACACCCCTCTGTGCCTCCTGACCTGACAGCCAGCG 1367

Qy   1289 CCACCACCTACCAGGGCAGTCTCTGTCCCCGGCAGGA----------------------- 1325
          ||  || |||||  ||     | | | ||| ||||||
Db   1368 CCGGCATCTACCGCGGACCCGTGTATGCCCTGCAGGACTCCACCGACAAAATCCCCATGA 1427

Qy   1326 ----------------------TGGGCCCAGCCCCAAGTTCCAGCTCACCA--------- 1354
                                ||||||| ||| || || || ||   ||
Db   1428 CCAACTCTCCTCTGCTGGACCCCTTACCCAGCCTTAAGGTCAAGGTCTACAGCTCCAGCA 1487

Qy   1355 --ATGGGCACCTGCTCAGCCCCCTGGGTGGCGGCCGCCACAC----------ACTGCACC 1402
            | |||| |  | |   ||| ||    |  | ||| | |           | ||
Db   1488 CCACGGGCTCTGGGCCAGGCCTGGCAGATGGGGCTGACCTGCTGGGGGTCTTGCCGCCTG 1547

Qy   1403 ACAGCTCTCCCACCTCTGAGGCCGAGGAGTTCGTCTCCCGCCTCTCCACCCAGAACTACT 1462
          || | || | | | |   |||  ||  ||    | || ||  |      |    |  | |
Db   1548 GCACATACCCTAGCGATTTCGCCCGGGACACCCACTTCCTGCACCTGCGCAGCGCCAGCC 1607

Qy   1463 TCCGCTC---------------CCTGCCCCGAGGCACCAGCAACATGACCTATGGGACCT 1507
          || | ||               |||||||||| | |  | | || |   |   || ||||
Db   1608 TCGGTTCCCAGCAGCTCTTGGGCCTGCCCCGAGACCCAGGGAGCAGCGTCAGCGGCACCT 1667

Qy   1508 TCAACTTCCTCGGGGGCCGGCTGATGATCCCTAATACAGGTATCAGCCTCCTCATCCCCC 1567
          |   || ||| || || |  || || |    ||||| | | |||||  |||||  |  |
Db   1668 TTGGCTGCCTGGGTGGGAGGCTCAGCATCCCCGGCACAGGGGTCAGCTTGCTGGTGCCCA 1727

Qy   1568 CAGATGCCATACCCCGAGGGAAGATCTATGAGATCTACCTCACGCTGCACAAGCCGGAAG 1627
          | ||||| |||| || | | ||  |   | |||| |||| | || || | |  |     |
Db   1728 ATGGAGCCATTCCCCAGGGCAAGTTCTACGAGATGTATCTACTCATCAACAAGGCAGAAA 1787

Qy   1628 ACGTGAGGTTGCCCCTAGCTGGCTGTCAGACCCTGCTGAGTCCCATCGTTAGCTGTGGAC 1687
```

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                    |  ||       ||   ||     |||||  |  ||||  |||    ||  | ||||||||
Db   1788 GTACCCTGCCGCTTTCAGAAGGGACCCAGACAGTATTGAGCCCCTCGGTGACCTGTGGAC 1847

Qy   1688 CCCCTGGCGTCCTGCTCACCCGGCCAGTCATCCTGGCTATGGACCACTGTGGGGAGCCCA 1747
          || | |||  |||||||     ||| || ||||||||  | ||  ||||||||  ||   ||
Db   1848 CCACAGGCCTCCTGCTGTGCCGCCCCGTCATCCTCACCATGCCCCACTGTGCCGAAGTCA 1907

Qy   1748 GCCCTGACAGCTGGAGCCTGCGCCTCAAAAAGCAGTCGTGCGAGGGCAGCTGGGAGGATG 1807
          |  |       |||||  |  |  |  ||||| |   ||| |   |  |||||  |||||||||| |
Db   1908 GTGCCCGTGACTGGATCTTTCAGCTCAAGACCCAGGCCCACCAGGGCCACTGGGAGGAGG 1967

Qy   1808 TGCTGCACCTGGGCGAGGAGGCGCCCTCCCACCTCTACTACTGCCAGCTGGAGGCCAGTG 1867
          ||  ||     |||||     ||||||  | |       |   | ||  |||||||||||||||||| ||||| |
Db   1968 TGGTGACCCTGGATGAGGAGACCCTGAACACACCCTGCTACTGCCAGCTGGAGCCCAGGG 2027

Qy   1868 CCTGCTACGTCTTCACCGAGCAGCTGGGCCGCTTTGCCCTGGTGGGAGAGGCCCTCAGCG 1927
          ||||   || ||  |     ||  ||||||||||  ||  |    | |   |||  ||| ||      |
Db   2028 CCTGTCACATCCTGCTGGACCAGCTGGGCACCTACGTGTTCACGGGCGAGTCCTATTCCC 2087

Qy   1928 TGGCTGCCGCCAAGCGCCTCAAGCTGCTTCTGTTTGCGCCGGTGGCCTGCACCTCCCTCG 1987
          |  || |  |||||||  |||  |||||  |   | || || || || |      |||||||||||||| |
Db   2088 GCTCAGCAGTCAAGCGGCTCCAGCTGGCCGTCTTCGCCCCCGCCCTCTGCACCTCCCTGG 2147

Qy   1988 AGTACAACATCCGGGTCTACTGCCTGCATGACACCCACGATGCACTCAAGGAGGTGGTGC 2047
          ||||||  |  ||||||||||||||||||||||||  | |||||| |  |   |||||  |||||||||| ||
Db   2148 AGTACAGCCTCCGGGTCTACTGCCTGGAGGACACGCCTGTAGCACTGAAGGAGGTGCTGG 2207

Qy   2048 AGCTGGAGAAGCAGCTGGGGGGGACAGCTGATCCAGGAGCCACGGGTCCTGCACTTCAAGG 2107
          ||||||||   |      ||||| |||  |   || |  |||||||         ||        |||||||
Db   2208 AGCTGGAGCGGACTCTGGGCGGATACTTGGTGGAGGAGCCGAAACCGCTAATGTTCAAGG 2267

Qy   2108 ACAGTTACCACAACCTGCGCCTATCCATCCACGATGTGCCCAGCTCCCTGTGGAAGAGTA 2167
          |||||||||||||||||||||||   |||  |||| ||   | |||     |||   |||| ||| |
Db   2268 ACAGTTACCACAACCTGCGCCTCTCCCTCCATGACCTCCCCCATGCCCATTGGAGGAGCA 2327

Qy   2168 AGCTCCTTGTCAGCTACCAGGAGATCCCCTTTTATCACATCTGGAATGGCACGCAGCGGT 2227
          ||||  || | ||  |||||||||||||||||||||||| |||||||| ||||  ||||| |||    |
Db   2328 AGCTGCTGGCCAAATACCAGGAGATCCCCTTCTATCACATTTGGAGTGGCAGCCAGAAGG 2387

Qy   2228 ACTTGCACTGCACCTTCACCCTGGAGCGTGTCAGCCCCAGCACTAGTGACCTGGCCTGCA 2287
          | | ||||||||| ||||||||||||||| |    ||||    | | | || || ||||||
Db   2388 CCCTCCACTGCACTTTCACCCTGGAGAGGCACAGCTTGGCCTCCACAGAGCTCACCTGCA 2447

Qy   2288 AGCTGTGGGTGTGGCAGGTGGAGGGCGACGGGCAGAGCTTCAGCATCAACTTCAACATCA 2347
          || | || |||  |||| || || || || |||| |||   |  |  |  ||   |
Db   2448 AGATCTGCGTGCGGCAAGTGGAAGGGGAGGGCCAGATATTCCAGCTGCATACCACTCTGG 2507

Qy   2348 CCAAG---GACACAAGGTTTGCTGAGCTGCTGGCTCTGGAGAGTGAAGCGGGGGTCCCAG 2404
          |  ||       |  |  |        ||     ||        |      ||        |||  |
Db   2508 CAGAGACACCTGCTGGCTCCCTGGACACTCTCTGCTCTGCCCCTGGCAGCACTGTCACCA 2567

Qy   2405 CCCTGGTGGGCCCCAGTGCCTTCAAGATCCCCTTCCTCATTCGGCAGAAGATAATTTCCA 2464
          ||| | ||||  ||     |||||||||||||||||||  |    ||| || ||||||||||          ||
Db   2568 CCCAGCTGGGACCTTATGCCTTCAAGATCCCACTGTCCATCCGCCAGAAGATATGCAACA 2627

Qy   2465 GCCTGGACCCACCCTGTAGGCGGGGTGCCGACTGGCGGACTCTGGCCCAGAAACTCCACC 2524
          ||||  ||  |  ||       |||||     |||||||||||    |  || ||||||  |||
```

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Db     2628  GCCTAGATGCCCCCAACTCACGGGGCAATGACTGGCGGATGTTAGCACAGAAGCTCTCTA 2687

Qy     2525  TGGACAGCCATCTCAGCTTCTTTGCCTCCAAGCCCAGCCCCACAGCCATGATCCTCAACC 2584
             |||| |  | || |   | ||||||| ||||  | ||||||||| |    ||||||| |||
Db     2688  TGGACCGGTACCTGAATTACTTTGCCACCAAAGCGAGCCCCACGGGTGTGATCCTGGACC 2747

Qy     2585  TGTGGGAGGCGCGGCACTTCCCCAACGGCAACCTCAGCCAGCTGGCTGCAGCAGTGGCTG 2644
             | |||||  ||  | |||       | | ||  ||||||  |  |||||   ||  ||| |
Db     2748  TCTGGGAAGCTCTGCAGCAGGACGATGGGGACCTCAACAGCCTGGCGAGTGCCTTGGAGG 2807

Qy     2645  GACTGGGCCAGCCAGACGCTGGCCTCTTCACAGTGTCGGAGGCTGAGTGCTGA 2697
             |||||  ||     ||         |      |  | || |  || ||||||
Db     2808  AGATGGGCAAGAGTGAGATGCTGGTGGCTGTGGCCACCGACGGGGACTGCTGA 2860


RESULT 14
US-10-218-779-1
; Sequence 1, Application US/10218779
; Publication No. US20040029222A1
; GENERAL INFORMATION:
;  APPLICANT: Edinger, Shlomit
;  APPLICANT:  MacDougall, John
;  APPLICANT:  Millet, Isabelle
;  APPLICANT:  Ellerman, Karen
;  APPLICANT:  Stone, David
;  APPLICANT:  Gerlach, Valerie
;  APPLICANT:  Grosse, William
;  APPLICANT:  Alsobrook II, John
;  APPLICANT:  Lepley, Denise
;  APPLICANT:  Rieger, Daniel
;  APPLICANT:  Burgess, Catherine
;  APPLICANT:  Casman, Stacie
;  APPLICANT:  Spytek, Kimberly
;  APPLICANT:  Boldog, Ferenc
;  APPLICANT:  Li, Li
;  APPLICANT:  Padigaru, Muralidhara
;  APPLICANT:  Mishra, Vishnu
;  APPLICANT:  Patturajan, Meera
;  APPLICANT:  Shenoy, Suresh
;  APPLICANT:  Rastelli, Luca
;  APPLICANT:  Tchernev, Velizar
;  APPLICANT:  Vernet, Corine
;  APPLICANT:  Zerhusen, Bryan
;  APPLICANT:  Malyankar, Uriel
;  APPLICANT:  Guo, Xiaojia
;  APPLICANT:  Miller, Charles
;  APPLICANT:  Gangolli, Esha
;  TITLE OF INVENTION: Proteins and Nucleic Acids Encoding Same
;  FILE REFERENCE: 21402-214
;  CURRENT APPLICATION NUMBER: US/10/218,779
;  CURRENT FILING DATE:  2002-08-14
;  PRIOR APPLICATION NUMBER: 60/253,834
;  PRIOR FILING DATE: 2000-11-29
;  PRIOR APPLICATION NUMBER: 60/250,-926
;  PRIOR FILING DATE: 2000-11-30
;  PRIOR APPLICATION NUMBER: 60/264,180
;  PRIOR FILING DATE: 2001-01-25
```

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  Query Match                34.7%;   Score 936.2;  DB 17;  Length 2860;
  Best Local Similarity      61.7%;   Pred. No. 2.8e-232;
  Matches 1662;  Conservative     0;  Mismatches  938;  Indels   93;  Gaps     7;


Qy        98 ACCCAGTGCCTGGTGCCAACCCGGACCTGCTTCCCCACTTCCTGGTGGAGCCCGAGGATG 157
             || | | || || |       | || | ||| ||| |||||||| |||||| |||| |
Db       168 ACTCCTTCCCGTCAGCGCCAGCAGAGCCGCTGCCCTACTTCCTGCAGGAGCCACAGGACG 227

Qy       158 TGTACATCGTCAAGAACAAGCCAGTGCTGCTTGTGTGCAAGGCCGTGCCCGCCACGCAGA 217
              ||||| || |||||||||||| |||  |||   |||     ||| | ||||||||| ||||
Db       228 CCTACATTGTGAAGAACAAGCCTGTGGAGCTCCGCTGCCGCGCCTTCCCCGCCACACAGA 287

Qy       218 TCTTCTTCAAGTGCAACGGGGAGTGGGTGCGCCAGGTGGACCACGTGATCGAGCGCAGCA 277
             ||| |||||||||||||||| |||||||| ||||| ||||||| | || ||
Db       288 TCTACTTCAAGTGCAACGGCGAGTGGGTCAGCCAGAACGACCACGTCACACAGGAAGGCC 347

Qy       278 CAGACGGGAGCAGTGGGCTGCCCACCATGGAGGTCCGCATTAATGTCTCAAGGCAGCAGG 337
              || | | || || |||| ||||| | || | || || |||||||||
Db       348 TGGATGAGGCCACCGGTCTGCGGGTGCGCGAGGTGCAGATCGAGGTGTCGCGGCAGCAGG 407

Qy       338 TCGAGAAGGTGTTCGGGCTGGAGGAATACTGGTGCCAGTGCGTGGCATGGAGCTCCTCGG 397
             | ||| || | | || ||||||||||| ||||||||||||||||||| ||||||||| |||
Db       408 TGGAGGAGCTCTTTGGGCTGGAGGATTACTGGTGCCAGTGCGTGGCCTGGAGCTCCGCGG 467

Qy       398 GCACCACCAAGAGTCAGAAGGCCTACATCCGCATAGCCAGATTGCGCAAGAACTTCGAGC 457
             ||||||||||||||||    |||||| ||||||| |||    |||||||||||||||| |
Db       468 GCACCACCAAGAGTCGCCGAGCCTACGTCCGCATCGCCTACCTGCGCAAGAACTTCGATC 527

Qy       458 AGGAGCCGCTGGCCAAGGAGGTGTCCCTGGAGCAGGGCATCGTGCTGCCCTGCCGTCCAC 517
             ||||||| |||| ||||||||||| |||||||||| || |   | | |||| ||||| || |
Db       528 AGGAGCCTCTGGGCAAGGAGGTGCCCCTGGACCATGAGGTTCTCCTGCAGTGCCGCCCGC 587

Qy       518 CGGAGGGCATCCCTCCAGCCGAGGTGGAGTGGCTCCGGAACGAGGACCTGGTGGACCCGT 577
             |||||| | | ||| |||||||||||| ||||||  |||  ||| ||||| | | |||||
Db       588 CGGAGGGGGGTGCCTGTGGCCGAGGTGGAATGGCTCAAGAATGAGGATGTCATCGACCCCA 647

Qy       578 CCCTGGACCCCAATGTATACATCACGCGGGAGCACAGCCTGGTGGTGCGACAGGCCCGCC 637
             ||| |||| |||| |     |||    || |||| ||| | | || |||||||||||
Db       648 CCCAGGACACCAACTTCCTGCTCACCATCGACCACAACCTCATCATCCGCCAGGCCCGCC 707

Qy       638 TTGCTGACACGGCCAACTACACCTGCGTGGCCAAGAACATCGTGGCACGTCGCCGCAGCG 697
             |  | ||||||| ||||||||| |||||||||||||||||||||||||||||     ||||| |||
Db       708 TGTCGGACACTGCCAACTATACCTGCGTGGCCAAGAACATCGTGGCCAAACGCCGGAGCA 767

Qy       698 CCTCCGCTGCTGTCATCGTCTACGTGAACGGTGGGTGGTCGACGTGGACCGAGTGGTCCG 757
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                           | |  |  | |     |  || |||||| ||||||||| || || ||||| |    ||| | |||||||||
Db     768 CCACTGCCACCGTCATCGTCTACGTGAATGGCGGCTGGTCCAGCTGGGCAGAGTGGTCAC 827

Qy     758 TCTGCAGCGCCAGCTGTGGGCGCGGCTGGCAGAAACGGAGCCGGAGCTGCACCAACCCGG 817
                           ||||  | · |  |||||||| || ||||||||||| || | |||||| |||||||||| |
Db     828 CCTGCTCCAACCGCTGTGGCCGAGGCTGGCAGAAGCGCACCCGGACCTGCACCAACCCCG 887

Qy     818 CGCCTCTCAACGGGGGCGCTTTCTGTGAGGGGCAGAATGTCCAGAAAACAGCCTGCGCCA 877
                           | | || |||||||| || || ||||| ||||| |||      ||||||| || |||||| |||
Db     888 CTCCACTCAACGGAGGGGCCTTCTGCGAGGGCCAGGCATTCCAGAAGACCGCCTGCACCA 947

Qy     878 CCCTGTGCCCAGTAGACGGCAGCTGGAGCCCGTGGAGCAAGTGGTCGGCCTGTGGGCTGG 937
                           || | |||||||| || ||     |||| ||||||||||||||| ||||| |    |
Db     948 CCATCTGCCCAGTCGATGGGGCGTGGACGGAGTGGAGCAAGTGGTCAGCCTGCAGCACTG 1007

Qy     938 ACTGCACCCACTGGCGGAGCCGTGAGTGCTCTGACCCAGCACCCCGCAACGGAGGGGAGG 997
                           | ||  |||||||||| ||||| ||||||    |  || ||||||  |||||||| |
Db    1008 AGTGTGCCCACTGGCGTAGCCGCGAGTGCATGGCGCCCCCACCCCAGAACGGAGGCCGTG 1067

Qy     998 AGTGCCAGGGCACTGACCTGGACACCCGCAACTGTACCAGTGACCTCTGTGTACACAGTG 1057
                           | |||    || ||      || ||| |      ||||| ||    || || || | ||      |
Db    1068 ACTGCAGCGGGACGCTGCTCGACTCTAAGAACTGCACAGATGGGCTGTGCATGCAACTGG 1127

Qy    1058 CTTCTGGCCCTGAGGACGTGGCCCTCTATGTGGGCCTCATCGCCGTGGCCGTCTGCCTGG 1117
                           |       || | | |     | | | | | ||| |   ||| |  | |||
Db    1128 AGGCCTCAGGGGATGCGGCGCTGTATGCGGGGCTCGTGGTGGCCATCTTCGTGGTCGTGG 1187

Qy    1118 TCCTGCTGCTGCTTGTCCTCATCCTCGTTTATTGCCGGAAGAAGGAGGGGCTGGACTCAG 1177
                           | || || ||         |  | | || ||  |||| ||        |   | ||| |||
Db    1188 CAATCCTCATGGCGGTGGGGGTGGTGGTGTACCGCCGCAACTGCCGTGACTTCGACACAG 1247

Qy    1178 ATGTGGCTGACTCGTCCATT---CTCACCTCAGGCTTCCAGCCCGTCAGCATCAAGCCCA 1234
                           | | | |||||||| ||   |   || ||     || ||||| ||||||| | | ||| |
Db    1248 ACATCACTGACTCATCTGCTGCCCTGACTGGTGGTTTCCACCCCGTCAACTTTAAGACGG 1307

Qy    1235 GCAAAGCAGACAACCCCCATCTGCT------CACCATCCAGCCGGACCTCAGCACCACCA 1288
                           |   |     ||||| || || ||      | | | || ||||| |   ||| |
Db    1308 CAAGGCCCAGTAACCCGCAGCTCCTACACCCCTCTGTGCCTCCTGACCTGACAGCCAGCG 1367

Qy    1289 CCACCACCTACCAGGGCAGTCTCTGTCCCCGGCAGGA---------------------- 1325
                           || || |||||| ||     | | | ||| ||||||
Db    1368 CCGGCATCTACCGCGGACCCGTGTATGCCCTGCAGGACTCCACCGACAAAATCCCCATGA 1427

Qy    1326 ----------------------TGGGCCCAGCCCCAAGTTCCAGCTCACCA--------- 1354
                                                 ||||||| ||| || || || ||   ||
Db    1428 CCAACTCTCCTCTGCTGGACCCCCTTACCCAGCCTTAAGGTCAAGGTCTACAGCTCCAGCA 1487

Qy    1355 --ATGGGCACCTGCTCAGCCCCCTGGGTGGCGGCCGCCACAC----------ACTGCACC 1402
                            | |||| |  |  ||| || |   | | ||| | |  |          | ||
Db    1488 CCACGGGCTCTGGGCCAGGCCTGGCAGATGGGGCTGACCTGCTGGGGGTCTTGCCGCCTG 1547

Qy    1403 ACAGCTCTCCCACCTCTGAGGCCGAGGAGTTCGTCTCCCGCCTCTCCACCCAGAACTACT 1462
                           ||  | || | | |   |||  |||  | |||| || | | |    |        | |
Db    1548 GCACATACCCTAGCGATTTCGCCCGGGACACCCACTTCCTGCACCTGCGCAGCGCCAGCC 1607

Qy    1463 TCCGCTC---------------CCTGCCCCGAGGCACCAGCAACATGACCTATGGGACCT 1507
                           || | ||               |||||||||||| || | | | | || |    |  || ||||
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Db   1608  TCGGTTCCCAGCAGCTCTTGGGCCTGCCCCGAGACCCAGGGAGCAGCGTCAGCGGCACCT  1667

Qy   1508  TCAACTTCCTCGGGGGCCGGCTGATGATCCCTAATACAGGTATCAGCCTCCTCATCCCCC  1567
           |  || ||| || ||  |||| |  |||||    ||||| ||||| | || | |||
Db   1668  TTGGCTGCCTGGGTGGGAGGCTCAGCATCCCCGGCACAGGGGTCAGCTTGCTGGTGCCCA  1727

Qy   1568  CAGATGCCATACCCCGAGGGAAGATCTATGAGATCTACCTCACGCTGCACAAGCCGGAAG  1627
            |  ||||| ||||   || |||  |||| |||||| || ||    |  ||||| | |||
Db   1728  ATGGAGCCATTCCCCAGGGCAAGTTCTACGAGATGTATCTACTCATCAACAAGGCAGAAA  1787

Qy   1628  ACGTGAGGTTGCCCCTAGCTGGCTGTCAGACCCTGCTGAGTCCCATCGTTAGCTGTGGAC  1687
            |  ||    ||  ||    ||||| |  |||| |||    || |  | ||||||||||
Db   1788  GTACCCTGCCGCTTTCAGAAGGGACCCAGACAGTATTGAGCCCCTCGGTGACCTGTGGAC  1847

Qy   1688  CCCCTGGCGTCCTGCTCACCCGGCCAGTCATCCTGGCTATGGACCACTGTGGGGAGCCCA  1747
           || | ||| |||||||   ||| || |||||||||| | |||| ||||||||| ||   ||
Db   1848  CCACAGGCCTCCTGCTGTGCCGCCCCGTCATCCTCACCATGCCCCACTGTGCCGAAGTCA  1907

Qy   1748  GCCCTGACAGCTGGAGCCTGCGCCTCAAAAAGCAGTCGTGCGAGGGCAGCTGGGAGGATG  1807
            |  |     ||||| | | |   ||||| |   ||| |   | ||||| |||||||||| |
Db   1908  GTGCCCGTGACTGGATCTTTCAGCTCAAGACCCAGGCCCACCAGGGCCACTGGGAGGAGG  1967

Qy   1808  TGCTGCACCTGGGCGAGGAGGCGCCCTCCCACCTCTACTACTGCCAGCTGGAGGCCAGTG  1867
           || ||  |||||   |||||| | |         | || |||||||||||||||| ||||| |
Db   1968  TGGTGACCCTGGATGAGGAGACCCTGAACACACCCTGCTACTGCCAGCTGGAGCCCAGGG  2027

Qy   1868  CCTGCTACGTCTTCACCGAGCAGCTGGGCCGCTTTGCCCTGGTGGGAGAGGCCCTCAGCG  1927
           ||||  || || |   || |||||||||| ||  |  |  ||| ||| || |        |
Db   2028  CCTGTCACATCCTGCTGGACCAGCTGGGCACCTACGTGTTCACGGGCGAGTCCTATTCCC  2087

Qy   1928  TGGCTGCCGCCAAGCGCCTCAAGCTGCTTCTGTTTGCGCCGGTGGCCTGCACCTCCCTCG  1987
            | || |  ||||||| ||| ||| |||||    | || || || |  ||||||||||||| |
Db   2088  GCTCAGCAGTCAAGCGGCTCCAGCTGGCCGTCTTCGCCCCCGCCCTCTGCACCTCCCTGG  2147

Qy   1988  AGTACAACATCCGGGTCTACTGCCTGCATGACACCCACGATGCACTCAAGGAGGTGGTGC  2047
           |||||| | |||||||||||||||||||||||| | ||||| |  |  ||||| |||||||| ||
Db   2148  AGTACAGCCTCCGGGTCTACTGCCTGGAGGACACGCCTGTAGCACTGAAGGAGGTGCTGG  2207

Qy   2048  AGCTGGAGAAGCAGCTGGGGGGGACAGCTGATCCAGGAGCCACGGGTCCTGCACTTCAAGG  2107
           |||||||| |   |||||  ||| ||| | || | |||||||       || |||||||| .
Db   2208  AGCTGGAGCGGACTCTGGGCGGATACTTGGTGGAGGAGCCGAAACCGCTAATGTTCAAGG  2267

Qy   2108  ACAGTTACCACAACCTGCGCCTATCCATCCACGATGTGCCCAGCTCCCTGTGGAAGAGTA  2167
           |||||||||||||||||||||| ||| |||| ||  |  | ||| |   |||| ||| |
Db   2268  ACAGTTACCACAACCTGCGCCTCTCCCTCCATGACCTCCCCCATGCCCATTGGAGGAGCA  2327
           · · ·
Qy   2168  AGCTCCTTGTCAGCTACCAGGAGATCCCCTTTTATCACATCTGGAATGGCACGCAGCGGT  2227
           |||| || | ||  |||||||||||||||||||||| |||||||| |||| |||||  ||| |
Db   2328  AGCTGCTGGCCAAATACCAGGAGATCCCCTTCTATCACATTTGGAGTGGCAGCCAGAAGG  2387

Qy   2228  ACTTGCACTGCACCTTCACCCTGGAGCGTGTCAGCCCCAGCACTAGTGACCTGGCCTGCA  2287
            | | |||||||| ||||||||||||| ||  ||||    |.| | || || |||||||
Db   2388  CCCTCCACTGCACTTTCACCCTGGAGAGGCACAGCTTGGCCTCCACAGAGCTCACCTGCA  2447

Qy   2288  AGCTGTGGGTGTGGCAGGTGGAGGGCGACGGGCAGAGCTTCAGCATCAACTTCAACATCA  2347
           || | || ||| |||| |||| || || || || ||||   ||    |  || |  ||
Db   2448  AGATCTGCGTGCGGCAAGTGGAAGGGGAGGGCCAGATATTCCAGCTGCATACCACTCTGG  2507
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Qy    2348 CCAAG---GACACAAGGTTTGCTGAGCTGCTGGCTCTGGAGAGTGAAGCGGGGGTCCCAG 2404
           |  ||      |  |  |     ||    ||        |   ||        ||| |
Db    2508 CAGAGACACCTGCTGGCTCCCTGGACACTCTCTGCTCTGCCCCTGGCAGCACTGTCACCA 2567

Qy    2405 CCCTGGTGGGCCCCAGTGCCTTCAAGATCCCCTTCCTCATTCGGCAGAAGATAATTTCCA 2464
           ||| | |||| ||    |||||||||||||||| |   ||| || |||||||||     ||
Db    2568 CCCAGCTGGGACCTTATGCCTTCAAGATCCCACTGTCCATCCGCCAGAAGATATGCAACA 2627

Qy    2465 GCCTGGACCCACCCTGTAGGCGGGGTGCCGACTGGCGGACTCTGGCCCAGAAACTCCACC 2524
           |||| || | |||     |||||     ||||||||| | || ||||| |||
Db    2628 GCCTAGATGCCCCCAACTCACGGGGCAATGACTGGCGGATGTTAGCACAGAAGCTCTCTA 2687

Qy    2525 TGGACAGCCATCTCAGCTTCTTTGCCTCCAAGCCCAGCCCCACAGCCATGATCCTCAACC 2584
           ||||| |   | || |  |||||||| ||||   | |||||||| |   |||||||  |||
Db    2688 TGGACCGGTACCTGAATTACTTTGCCACCAAAGCGAGCCCCACGGGTGTGATCCTGGACC 2747

Qy    2585 TGTGGGAGGCGCGGCACTTCCCCAACGGCAACCTCAGCCAGCTGGCTGCAGCAGTGGCTG 2644
           | ||||| || | |||    | |  ||  ||||| |   ||||| |   ||  ||| |
Db    2748 TCTGGGAAGCTCTGCAGCAGGACGATGGGGACCTCAACAGCCTGGCGAGTGCCTTGGAGG 2807

Qy    2645 GACTGGGCCAGCCAGACGCTGGCCTCTTCACAGTGTCGGAGGCTGAGTGCTGA 2697
           ||||| ||   |.|        |       |   | || |  || ||||||
Db    2808 AGATGGGCAAGAGTGAGATGCTGGTGGCTGTGGCCACCGACGGGGACTGCTGA 2860
```

RESULT 15
US-10-218-779-3
; Sequence 3, Application US/10218779
; Publication No. US20040029222A1
; GENERAL INFORMATION:
; APPLICANT: Edinger, Shlomit
; APPLICANT: MacDougall, John
; APPLICANT: Millet, Isabelle
; APPLICANT: Ellerman, Karen
; APPLICANT: Stone, David
; APPLICANT: Gerlach, Valerie
; APPLICANT: Grosse, William
; APPLICANT: Alsobrook II, John
; APPLICANT: Lepley, Denise
; APPLICANT: Rieger, Daniel
; APPLICANT: Burgess, Catherine
; APPLICANT: Casman, Stacie
; APPLICANT: Spytek, Kimberly
; APPLICANT: Boldog, Ferenc
; APPLICANT: Li, Li
; APPLICANT: Padigaru, Muralidhara
; APPLICANT: Mishra, Vishnu
; APPLICANT: Patturajan, Meera
; APPLICANT: Shenoy, Suresh
; APPLICANT: Rastelli, Luca
; APPLICANT: Tchernev, Velizar
; APPLICANT: Vernet, Corine
; APPLICANT: Zerhusen, Bryan
; APPLICANT: Malyankar, Uriel
; APPLICANT: Guo, Xiaojia
; APPLICANT: Miller, Charles

```
    Query Match               34.7%;   Score 936.2;  DB 17;   Length 2860;
    Best Local Similarity     61.7%;   Pred. No. 2.8e-232;
    Matches 1662;  Conservative    0;  Mismatches  938;  Indels  93;  Gaps    7;

Qy       98 ACCCAGTGCCTGGTGCCAACCCGGACCTGCTTCCCCACTTCCTGGTGGAGCCCGAGGATG 157
            || |  | ||   ||       | || | ||| ||| |||||||| |||||| |||| |
Db      168 ACTCCTTCCCGTCAGCGCCAGCAGAGCCGCTGCCCTACTTCCTGCAGGAGCCACAGGACG 227

Qy      158 TGTACATCGTCAAGAACAAGCCAGTGCTGCTTGTGTGCAAGGCCGTGCCCGCCACGCAGA 217
              ||||| ||  ||||||||||||| ||| |||| ||| | |||  ||| | ||||||||| ||||
Db      228 CCTACATTGTGAAGAACAAGCCTGTGGAGCTTCGCTGCCGCGCCTTCCCCGCCACACAGA 287

Qy      218 TCTTCTTCAAGTGCAACGGGGAGTGGGTGCGCCAGGTGGACCACGTGATCGAGCGCAGCA 277
            ||| |||||||||||||||| |||||||| ||||||||| |||||     |||||||| |   ||        ||
Db      288 TCTACTTCAAGTGCAACGGCGAGTGGGTCAGCCAGAACGACCACGTCACACAGGAAGGCC 347

Qy      278 CAGACGGGAGCAGTGGGCTGCCCACCATGGAGGTCCGCATTAATGTCTCAAGGCAGCAGG 337
               || | |   ||  || ||||         |·|||| | | || | | ||·|| ·| ·|·||||||·
Db      348 TGGATGAGGCCACCGGCCTGCGGGTGCGCGAGGTGCAGATCGAGGTGTCGCGGCAGCAGG 407

Qy      338 TCGAGAAGGTGTTCGGGCTGGAGGAATACTGGTGCCAGTGCGTGGCATGGAGCTCCTCGG 397
            | ||| || | ||  ||||||||||| |||||||||||||||||||||||||| |||||||||| | |
Db      408 TGGAGGAGCTCTTTGGGCTGGAGGATTACTGGTGCCAGTGCGTGGCCTGGAGCTCCGCAG 467

Qy      398 GCACCACCAAGAGTCAGAAGGCCTACATCCGCATAGCCAGATTGCGCAAGAACTTCGAGC 457
            |||||||||||||||      ||||||| ||||||| |||      |||||||||||||||| |
Db      468 GCACCACCAAGAGTCGCCGAGCCTACGTCCGCATCGCCTACCTGCGCAAGAACTTCGATC 527

Qy      458 AGGAGCCGCTGGCCAAGGAGGTGTCCCTGGAGCAGGGCATCGTGCTGCCCTGCCGTCCAC 517
            ||||||| |||| ||||||||||| |||||||| || |    |  | |||| ||||| || |
Db      528 AGGAGCCTCTGGGCAAGGAGGTGCCCCTGGACCATGAGGTTCTCCTGCAGTGCCGCCCGC 587

Qy      518 CGGAGGGCATCCCTCCAGCCGAGGTGGAGTGGCTCCGGAACGAGGACCTGGTGGACCCGT 577
               |||||||| | |||    ||||||||||| |||||| ||| |||||  |  | |||||
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Db    588  CGGAGGGGGTGCCTGTGGCCGAGGTGGAATGGCTCAAGAATGAGGATGTCATCGACCCCA  647

Qy    578  CCCTGGACCCCAATGTATACATCACGCGGGAGCACAGCCTGGTGGTGCGACAGGCCCGCC  637
           ||| |||| ||||  |      ||||    || |||| |||  |  | || |||||||||
Db    648  CCCAGGACACCAACTTCCTGCTCACCATCGACCACAACCTCATCATCCGCCAGGCCCGCC  707

Qy    638  TTGCTGACACGGCCAACTACACCTGCGTGGCCAAGAACATCGTGGCACGTCGCCGCAGCG  697
           |  |  |||||| ||||||||| |||||||||||||||||||||||||||||   ||||| |||
Db    708  TGTCGGACACTGCCAACTATACCTGCGTGGCCAAGAACATCGTGGCCAAACGCCGGAGCA  767

Qy    698  CCTCCGCTGCTGTCATCGTCTACGTGAACGGTGGGTGGTCGACGTGGACCGAGTGGTCCG  757
           || | ||  | |||||||||||||||||||| || || ||||| |  ||| | ||||||||
Db    768  CCACTGCCACCGTCATCGTCTACGTGAATGGCGGCTGGTCCAGCTGGGCAGAGTGGTCAC  827

Qy    758  TCTGCAGCGCCAGCTGTGGGCGCGGCTGGCAGAAACGGAGCCGGAGCTGCACCAACCCGG  817
           ||||   |  | |||||||| ||  |||||||||||  || |  |||||| |||||||||||| |
Db    828  CCTGCTCCAACCGCTGTGGCCGAGGCTGGCAGAAGCGCACCCGGACCTGCACCAACCCCG  887

Qy    818  CGCCTCTCAACGGGGGCGCTTTCTGTGAGGGGCAGAATGTCCAGAAAACAGCCTGCGCCA  877
           | || |||||||| ||  || || ||  |||||  |||||  |||   ||||||| || ||||||| |||
Db    888  CTCCACTCAACGGAGGGGCCTTCTGCGAGGGCCAGGCATTCCAGAAGACCGCCTGCACCA  947

Qy    878  CCCTGTGCCCAGTAGACGGCAGCTGGAGCCCGTGGAGCAAGTGGTCGGCCTGTGGGCTGG  937
           || | |||||||| || ||     |||| |||||||||||||||||| ||||| |    |
Db    948  CCATCTGCCCAGTCGATGGGGCGTGGACGGAGTGGAGCAAGTGGTCAGCCTGCAGCACTG  1007

Qy    938  ACTGCACCCACTGGCGGAGCCGTGAGTGCTCTGACCCAGCACCCCGCAACGGAGGGGAGG  997
           | ||  |||||||||| ||||| ||||||   |  || |||||| |||||||| |
Db    1008 AGTGTGCCCACTGGCGTAGCCGCGAGTGCATGGCGCCCCCACCCCAGAACGGAGGCCGTG  1067

Qy    998  AGTGCCAGGGCACTGACCTGGACACCCGCAACTGTACCAGTGACCTCTGTGTACACAGTG  1057
           | |||   || || ||  || |||  | |||||| ||  ||  || || || | ||
Db    1068 ACTGCAGCGGGACGCTGCTCGACTCTAAGAACTGCACAGATGGGCTGTGCATGCAACTGG  1127

Qy    1058 CTTCTGGCCCTGAGGACGTGGCCCTCTATGTGGGCCTCATCGCCGTGGCCGTCTGCCTGG  1117
           |          || |·| ||          | |  | |  | ||| |   ||    | |||
Db    1128 AGGCCTCAGGGGATGCGGCGCTGTATGCGGGGCTCGTGGTGGCCATCTTCGTGGTCGTGG  1187

Qy    1118 TCCTGCTGCTGCTTGTCCTCATCCTCGTTTATTGCCGGAAGAAGGAGGGGCTGGACTCAG  1177
            | ||  ||   ||     | | || || |||| ||      |   | ||| |||
Db    1188 CAATCCTCATGGCGGTGGGGGTGGTGGTGTACCGCCGCAACTGCCGTGACTTCGACACAG  1247

Qy    1178 ATGTGGCTGACTCGTCCATT---CTCACCTCAGGCTTCCAGCCCGTCAGCATCAAGCCCA  1234
           | |  |||||||| ||  |     || ||  || ||||| |||||||| | | ||| |
Db    1248 ACATCACTGACTCATCTGCTGCCCTGACTGGTGGTTTCCACCCCGTCAACTTTAAGACGG  1307

Qy    1235 GCAAAGCAGACAACCCCCATCTGCT------CACCATCCAGCCGGACCTCAGCACCACCA  1288
           |    | |     ||||| || || ||        | | | | || || ||||| |   ||| |
Db    1308 CAAGGCCCAGTAACCCGCAGCTCCTACACCCCTCTGTGCCTCCTGACCTGACAGCCAGCG  1367

Qy    1289 CCACCACCTACCAGGGCAGTCTCTGTCCCCGGCAGGA----------------------- 1325
           ||   || ||||||  ||       |  | ||| ||||||
Db    1368 CCGGCATCTACCGCGGACCCGTGTATGCCCTGCAGGACTCCACCGACAAAATCCCCATGA  1427

Qy    1326 -----------------------TGGGCCCAGCCCCAAGTTCCAGCTCACCA--------- 1354
                                  ||||||| ||| || || || |   ||
Db    1428 CCAACTCTCCTCTGCTGGACCCCTTACCCAGCCTTAAGGTCAAGGTCTACAGCTCCAGCA  1487
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Qy   1355 --ATGGGCACCTGCTCAGCCCCCTGGGTGGCGGCCGCCACAC----------ACTGCACC 1402
            | ||||| |  |   ||| ||      |  | |||| |  |   |          | ||
Db   1488 CCACGGGCTCTGGGCCAGGCCTGGCAGATGGGGCTGACCTGCTGGGGGTCTTGCCGCCTG 1547

Qy   1403 ACAGCTCTCCCACCTCTGAGGCCGAGGAGTTCGTCTCCCGCCTCTCCACCCAGAACTACT 1462
            ||  |  || | |  |     ||| |||  |  || ||  ||    |        |  |
Db   1548 GCACATACCCTAGCGATTTCGCCCGGGACACCCACTTCCTGCACCTGCGCAGCGCCAGCC 1607

Qy   1463 TCCGCTC---------------CCTGCCCCGAGGCACCAGCAACATGACCTATGGGACCT 1507
            || | ||               ||||||||||| | |  | | ||      |  || ||||
Db   1608 TCGGTTCCCAGCAGCTCTTGGGCCTGCCCCGAGACCCAGGGAGCAGCGTCAGCGGCACCT 1667

Qy   1508 TCAACTTCCTCGGGGGCCGGCTGATGATCCCTAATACAGGTATCAGCCTCCTCATCCCCC 1567
            |   || ||| || || ||  |||| |  |||||     |||||  |||||| | ||  |||
Db   1668 TTGGCTGCCTGGGTGGGAGGCTCAGCATCCCCGGCACAGGGGTCAGCTTGCTGGTGCCCA 1727

Qy   1568 CAGATGCCATACCCCGAGGGAAGATCTATGAGATCTACCTCACGCTGCACAAGCCGGAAG 1627
            |  ||||| ||||  || |||  ||| ||||  ||||| ||||| || ||     |  ||| | |||
Db   1728 ATGGAGCCATTCCCCAGGGCAAGTTCTACGAGATGTATCTACTCATCAACAAGGCAGAAA 1787

Qy   1628 ACGTGAGGTTGCCCCTAGCTGGCTGTCAGACCCTGCTGAGTCCCATCGTTAGCTGTGGAC 1687
             |  ||      ||  ||  ||     ||||| |  |||| |||  || | |||||||||
Db   1788 GTACCCTGCCGCTTTCAGAAGGGACCCAGACAGTATTGAGCCCCTCGGTGACCTGTGGAC 1847

Qy   1688 CCCCTGGCGTCCTGCTCACCCGGCCAGTCATCCTGGCTATGGACCACTGTGGGGAGCCCA 1747
            || | ||| |||||||    ||| || |||||||||| | |||| ||||||||  ||  ||
Db   1848 CCACAGGCCTCCTGCTGTGCCGCCCCGTCATCCTCACCATGCCCCACTGTGCCGAAGTCA 1907

Qy   1748 GCCCTGACAGCTGGAGCCTGCGCCTCAAAAAGCAGTCGTGCGAGGGCAGCTGGGAGGATG 1807
            |  |     ||||| | |  |  |||||| |  | ||| |    | |||||  |||||||||| |
Db   1908 GTGCCCGTGACTGGATCTTTCAGCTCAAGACCCAGGCCCACCAGGGCCACTGGGAGGAGG 1967

Qy   1808 TGCTGCACCTGGGCGAGGAGGCGCCCTCCCACCTCTACTACTGCCAGCTGGAGGCCAGTG 1867
            || ||  |||||| |||||| | |    |      | || |||||||||||||||||| |||| |
Db   1968 TGGTGACCCTGGATGAGGAGACCCTGAACACACCCTGCTACTGCCAGCTGGAGCCCAGGG 2027

Qy   1868 CCTGCTACGTCTTCACCGAGCAGCTGGGCCGCTTTGCCCTGGTGGGAGAGGCCCTCAGCG 1927
            ||||  || || |   || || |||||||||| || |  |   ||| ||| || |
Db   2028 CCTGTCACATCCTGCTGGACCAGCTGGGCACCTACGTGTTCACGGGCGAGTCCTATTCCC 2087

Qy   1928 TGGCTGCCGCCAAGCGCCTCAAGCTGCTTCTGTTTGCGCCGGTGGCCTGCACCTCCCTCG 1987
             | || | |||||| ||| ||| |||||  | || || ||  | |||||||||||| |
Db   2088 GCTCAGCAGTCAAGCGGCTCCAGCTGGCCGTCTTCGCCCCCGCCCTCTGCACCTCCCTGG 2147

Qy   1988 AGTACAACATCCGGGTCTACTGCCTGCATGACACCCACGATGCACTCAAGGAGGTGGTGC 2047 --
            ||||||  | ||||||||||||||||||||||| | |||||| |  | ||||| |||||||||| ||
Db   2148 AGTACAGCCTCCGGGTCTACTGCCTGGAGGACACGCCTGTAGCACTGAAGGAGGTGCTGG 2207

Qy   2048 AGCTGGAGAAGCAGCTGGGGGGACAGCTGATCCAGGAGCCACGGGTCCTGCACTTCAAGG 2107
            ||||||||| |     ||||| ||| | || |   |||||||        || ||||||||
Db   2208 AGCTGGAGCGGACTCTGGGCGGATACTTGGTGGAGGAGCCGAAACCGCTAATGTTCAAGG 2267

Qy   2108 ACAGTTACCACAACCTGCGCCTATCCATCCACGATGTGCCCAGCTCCCTGTGGAAGAGTA 2167
            ||||||||||||||||||||||| |  ||| |||| | | |||        ||| |||| ||| |
Db   2268 ACAGTTACCACAACCTGCGCCTCTCCCTCCATGACCTCCCCCATGCCCATTGGAGGAGCA 2327
```

```
Qy    2168 AGCTCCTTGTCAGCTACCAGGAGATCCCCTTTTATCACATCTGGAATGGCACGCAGCGGT 2227
            -||||  ||  |  ||   ||||||||||||||||||| ||||||||| |||| ||||||. |||   |
Db    2328 AGCTGCTGGCCAAATACCAGGAGATCCCCTTCTATCACATTTGGAGTGGCAGCCAGAAGG 2387

Qy    2228 ACTTGCACTGCACCTTCACCCTGGAGCGTGTCAGCCCCAGCACTAGTGACCTGGCCTGCA 2287
            |  |  |||||||||  |||||||||||||  |    ||||     |  |   || ||   ||||||
Db    2388 CCCTCCACTGCACTTTCACCCTGGAGAGGCACAGCTTGGCCTCCACAGAGCTCACCTGCA 2447

Qy    2288 AGCTGTGGGTGTGGCAGGTGGAGGGCGACGGGCAGAGCTTCAGCATCAACTTCAACATCA 2347
            || | || |||  |||| |||||  |||||| ||  || ||  |||||   |  |    || |
Db    2448 AGATCTGCGTGCGGCAAGTGGAAGGGGAGGGCCAGATATTCCAGCTGCATACCACTCTGG 2507

Qy    2348 CCAAG---GACACAAGGTTTGCTGAGCTGCTGGCTCTGGAGAGTGAAGCGGGGGTCCCAG 2404
            |  ||       |  |  |   |·|    ||    |     |    ||    ||| |
Db    2508 CAGAGACACCTGCTGGCTCCCTGGACACTCTCTGCTCTGCCCCTGGCAGCACTGTCACCA 2567

Qy    2405 CCCTGGTGGGCCCCAGTGCCTTCAAGATCCCCTTCCTCATTCGGCAGAAGATAATTTCCA 2464
            .||| | ||||  ||   ||||||||||||||||||||  |    ||| ||  ||||||||||    ||
Db    2568 CCCAGCTGGGACCTTATGCCTTCAAGATCCCACTGTCCATCCGCCAGAAGATATGCAACA 2627

Qy    2465 GCCTGGACCCACCCTGTAGGCGGGGTGCCGACTGGCGGACTCTGGCCCAGAAACTCCACC 2524
            ||||  ||   |  |||     |||||    ||||||||||  |  || |||||| |||
Db    2628 GCCTAGATGCCCCCAACTCACGGGGCAATGACTGGCGGATGTTAGCACAGAAGCTCTCTA 2687

Qy    2525 TGGACAGCCATCTCAGCTTCTTTGCCTCCAAGCCCAGCCCCACAGCCATGATCCTCAACC 2584
            |||||  |    | ||  |    | ||||||| ||||  | ||||||||| |    |||||||  |||
Db    2688 TGGACCGGTACCTGAATTACTTTGCCACCAAAGCGAGCCCCACGGGTGTGATCCTGGACC 2747

Qy    2585 TGTGGGAGGCGCGGCACTTCCCCAACGGCAACCTCAGCCAGCTGGCTGCAGCAGTGGCTG 2644
            | |||||| || |  |||      |  | || |||||| |  |||||| ·  ||  |||  |
Db    2748 TCTGGGAAGCTCTGCAGCAGGACGATGGGGACCTCAACAGCCTGGCGAGTGCCTTGGAGG 2807

Qy    2645 GACTGGGCCAGCCAGACGCTGGCCTCTTCACAGTGTCGGAGGCTGAGTGCTGA 2697
            ||||| ||   ||       |        |     |  | ||  |  || ||||||
Db    2808 AGATGGGCAAGAGTGAGATGCTGGTGGCTGTGGCCACCGACGGGGACTGCTGA 2860


Search completed: March  6, 2005, 17:04:14
Job time : 1484.63 secs
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